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# *Streptococcus pneumoniae* Antigens and Vaccines

## *Field of the Invention*

The present invention relates to novel *Streptococcus pneumoniae* antigens for the detection of *Streptococcus* and for the prevention or attenuation of disease caused by *Streptococcus*. The invention further relates to isolated nucleic acid molecules encoding antigenic polypeptides of *S. pneumoniae*. Antigenic polypeptides are also provided, as are vectors, host cells and recombinant methods for producing the same. The invention additionally relates to diagnostic methods for detecting *Streptococcus* gene expression.

## *Background of the Invention*

*Streptococcus pneumoniae* has been one of the most extensively studied microorganisms since its first isolation in 1881. It was the object of many investigations that led to important scientific discoveries. In 1928, Griffith observed that when heat-killed encapsulated pneumococci and live strains constitutively lacking any capsule were concomitantly injected into mice, the nonencapsulated could be converted into encapsulated pneumococci with the same capsular type as the heat-killed strain. Years later, the nature of this "transforming principle," or carrier of genetic information, was shown to be DNA. (Avery, O.T., et al., *J. Exp. Med.*, 79:137-157 (1944)).

In spite of the vast number of publications on *S. pneumoniae* many questions about its virulence are still unanswered, and this pathogen remains a major causative agent of serious human disease, especially community-acquired pneumonia. (Johnston, R.B., et al., *Rev. Infect. Dis.* 13(Suppl. 6):S509-517 (1991)). In addition, in developing countries, the pneumococcus is responsible for the death of a large number of children under the age of 5 years from pneumococcal pneumonia. The incidence of pneumococcal disease is highest in infants under 2 years of age and in people over 60 years of age. Pneumococci are the second most frequent cause (after *Haemophilus influenzae* type b) of bacterial meningitis and otitis media in children. With the recent introduction of conjugate vaccines for *H. influenzae* type b, pneumococcal meningitis is likely to become increasingly prominent. *S. pneumoniae* is the most important etiologic agent of community-acquired pneumonia in adults and is the second most common cause of bacterial meningitis behind *Neisseria meningitidis*.

The antibiotic generally prescribed to treat *S. pneumoniae* is benzylpenicillin, although resistance to this and to other antibiotics is found occasionally. Pneumococcal resistance to penicillin results from mutations in its

penicillin-binding proteins. In uncomplicated pneumococcal pneumonia caused by a sensitive strain, treatment with penicillin is usually successful unless started too late. Erythromycin or clindamycin can be used to treat pneumonia in patients hypersensitive to penicillin, but resistant strains to these drugs exist. Broad spectrum antibiotics (e.g., the tetracyclines) may also be effective, although tetracycline-resistant strains are not rare. In spite of the availability of antibiotics, the mortality of pneumococcal bacteremia in the last four decades has remained stable between 25 and 29%. (Gillespie, S.H., *et al.*, *J. Med. Microbiol.* 28:237-248 (1989).

*S. pneumoniae* is carried in the upper respiratory tract by many healthy individuals. It has been suggested that attachment of pneumococci is mediated by a disaccharide receptor on fibronectin, present on human pharyngeal epithelial cells. (Anderson, B.J., *et al.*, *J. Immunol.* 142:2464-2468 (1989). The mechanisms by which pneumococci translocate from the nasopharynx to the lung, thereby causing pneumonia, or migrate to the blood, giving rise to bacteremia or septicemia, are poorly understood. (Johnston, R.B., *et al.*, *Rev. Infect. Dis.* 13(Suppl. 6):S509-517 (1991).

Various proteins have been suggested to be involved in the pathogenicity of *S. pneumoniae*, however, only a few of them have actually been confirmed as virulence factors. Pneumococci produce an IgA1 protease that might interfere with host defense at mucosal surfaces. (Kornfield, S.J., *et al.*, *Rev. Inf. Dis.* 3:521-534 (1981). *S. pneumoniae* also produces neuraminidase, an enzyme that may facilitate attachment to epithelial cells by cleaving sialic acid from the host glycolipids and gangliosides. Partially purified neuraminidase was observed to induce meningitis-like symptoms in mice; however, the reliability of this finding has been questioned because the neuraminidase preparations used were probably contaminated with cell wall products. Other pneumococcal proteins besides neuraminidase are involved in the adhesion of pneumococci to epithelial and endothelial cells. These pneumococcal proteins have as yet not been identified. Recently, Cundell *et al.*, reported that peptide permeases can modulate pneumococcal adherence to epithelial and endothelial cells. It was, however, unclear whether these permeases function directly as adhesions or whether they enhance adherence by modulating the expression of pneumococcal adhesions. (DeVelasco, E.A., *et al.*, *Micro. Rev.* 59:591-603 (1995). A better understanding of the virulence factors determining its pathogenicity will need to be developed to cope with the devastating effects of pneumococcal disease in humans.

Ironically, despite the prominent role of *S. pneumoniae* in the discovery of DNA, little is known about the molecular genetics of the organism. The *S. pneumoniae* genome consists of one circular, covalently closed, double-stranded DNA and a collection of so-called variable accessory elements, such as prophages, plasmids, transposons and the like. Most physical characteristics and almost all of the genes of *S. pneumoniae* are unknown. Among the few that have been identified, most have not been physically mapped or characterized in detail. Only a few genes of this organism have been sequenced. (See, for instance current versions of GENBANK and other nucleic acid databases, and references that relate to the genome of *S. pneumoniae* such as those set out elsewhere herein.) Identification of *in vivo*-expressed, and broadly protective, antigens of *S. pneumoniae* has remained elusive.

#### *Summary of the Invention*

The present invention provides isolated nucleic acid molecules comprising polynucleotides encoding the *S. pneumoniae* polypeptides described in Table 1 and having the amino acid sequences shown as SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, and so on through SEQ ID NO:226. Thus, one aspect of the invention provides isolated nucleic acid molecules comprising polynucleotides having a nucleotide sequence selected from the group consisting of: (a) a nucleotide sequence encoding any of the amino acid sequences of the polypeptides shown in Table 1; and (b) a nucleotide sequence complementary to any of the nucleotide sequences in (a).

Further embodiments of the invention include isolated nucleic acid molecules that comprise a polynucleotide having a nucleotide sequence at least 90% identical, and more preferably at least 95%, 96%, 97%, 98% or 99% identical, to any of the nucleotide sequences in (a) or (b) above, or a polynucleotide which hybridizes under stringent hybridization conditions to a polynucleotide in (a) or (b) above. This polynucleotide which hybridizes does not hybridize under stringent hybridization conditions to a polynucleotide having a nucleotide sequence consisting of only A residues or of only T residues. Additional nucleic acid embodiments of the invention relate to isolated nucleic acid molecules comprising polynucleotides which encode the amino acid sequences of epitope-bearing portions of an *S. pneumoniae* polypeptide having an amino acid sequence in (a) above.

The present invention also relates to recombinant vectors, which include the isolated nucleic acid molecules of the present invention, and to host cells containing the recombinant vectors, as well as to methods of making such

vectors and host cells and for using these vectors for the production of *S. pneumoniae* polypeptides or peptides by recombinant techniques.

The invention further provides isolated *S. pneumoniae* polypeptides having an amino acid sequence selected from the group consisting of an amino acid sequence of any of the polypeptides described in Table 1.

The polypeptides of the present invention also include polypeptides having an amino acid sequence with at least 70% similarity, and more preferably at least 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% similarity to those described in Table 1, as well as polypeptides having an amino acid sequence at least 70% identical, more preferably at least 75% identical, and still more preferably 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% identical to those above; as well as isolated nucleic acid molecules encoding such polypeptides.

The present invention further provides a vaccine, preferably a multi-component vaccine comprising one or more of the *S. pneumoniae* polynucleotides or polypeptides described in Table 1, or fragments thereof, together with a pharmaceutically acceptable diluent, carrier, or excipient, wherein the *S. pneumoniae* polypeptide(s) are present in an amount effective to elicit an immune response to members of the *Streptococcus* genus in an animal. The *S. pneumoniae* polypeptides of the present invention may further be combined with one or more immunogens of one or more other streptococcal or non-streptococcal organisms to produce a multi-component vaccine intended to elicit an immunological response against members of the *Streptococcus* genus and, optionally, one or more non-streptococcal organisms.

The vaccines of the present invention can be administered in a DNA form, e.g., "naked" DNA, wherein the DNA encodes one or more streptococcal polypeptides and, optionally, one or more polypeptides of a non-streptococcal organism. The DNA encoding one or more polypeptides may be constructed such that these polypeptides are expressed fusion proteins.

The vaccines of the present invention may also be administered as a component of a genetically engineered organism. Thus, a genetically engineered organism which expresses one or more *S. pneumoniae* polypeptides may be administered to an animal. For example, such a genetically engineered organism may contain one or more *S. pneumoniae* polypeptides of the present invention intracellularly, on its cell surface, or in its periplasmic space. Further, such a genetically engineered organism may secrete one or more *S. pneumoniae* polypeptides.



The vaccines of the present invention may be co-administered to an animal with an immune system modulator (*e.g.*, CD86 and GM-CSF).

The invention also provides a method of inducing an immunological response in an animal to one or more members of the *Streptococcus* genus, preferably one or more isolates of the *S. pneumoniae* genus, comprising administering to the animal a vaccine as described above.

The invention further provides a method of inducing a protective immune response in an animal, sufficient to prevent or attenuate an infection by members of the *Streptococcus* genus, preferably at least *S. pneumoniae*, comprising administering to the animal a composition comprising one or more of the polynucleotides or polypeptides described in Table 1, or fragments thereof. Further, these polypeptides, or fragments thereof, may be conjugated to another immunogen and/or administered in admixture with an adjuvant.

The invention further relates to antibodies elicited in an animal by the administration of one or more *S. pneumoniae* polypeptides of the present invention and to methods for producing such antibodies.

The invention also provides diagnostic methods for detecting the expression of genes of members of the *Streptococcus* genus in an animal. One such method involves assaying for the expression of a gene encoding *S. pneumoniae* peptides in a sample from an animal. This expression may be assayed either directly (*e.g.*, by assaying polypeptide levels using antibodies elicited in response to amino acid sequences described in Table 1) or indirectly (*e.g.*, by assaying for antibodies having specificity for amino acid sequences described in Table 1). An example of such a method involves the use of the polymerase chain reaction (PCR) to amplify and detect *Streptococcus* nucleic acid sequences.

The present invention also relates to nucleic acid probes having all or part of a nucleotide sequence described in Table 1 (shown as SEQ ID NO:1, SEQ ID NO:3, SEQ ID NO:5, and so on through SEQ ID NO:225) which are capable of hybridizing under stringent conditions to *Streptococcus* nucleic acids. The invention further relates to a method of detecting one or more *Streptococcus* nucleic acids in a biological sample obtained from an animal, said one or more nucleic acids encoding *Streptococcus* polypeptides, comprising: (a) contacting the sample with one or more of the above-described nucleic acid probes, under conditions such that hybridization occurs, and (b) detecting hybridization of said one or more probes to the *Streptococcus* nucleic acid present in the biological sample.

The invention also includes immunoassays, including an immunoassay for detecting *Streptococcus*, preferably at least isolates of the *S. pneumoniae* genus, comprising incubation of a sample (which is suspected of being infected with *Streptococcus*) with a probe antibody directed against an antigen/epitope of *S. pneumoniae*, to be detected under conditions allowing the formation of an antigen-antibody complex; and detecting the antigen-antibody complex which contains the probe antibody. An immunoassay for the detection of antibodies which are directed against a *Streptococcus* antigen comprising the incubation of a sample (containing antibodies from a mammal suspected of being infected with *Streptococcus*) with a probe polypeptide including an epitope of *S. pneumoniae*, under conditions that allow the formation of antigen-antibody complexes which contain the probe epitope containing antigen.

Some aspects of the invention pertaining to kits are those for: investigating samples for the presence of polynucleotides derived from *Streptococcus* which comprise a polynucleotide probe including a nucleotide sequence selected from Table 1 or a fragment thereof of approximately 15 or more nucleotides, in an appropriate container; analyzing the samples for the presence of antibodies directed against a *Streptococcus* antigen made up of a polypeptide which contains a *S. pneumoniae* epitope present in the polypeptide, in a suitable container; and analyzing samples for the presence of *Streptococcus* antigens made up of an anti-*S. pneumoniae* antibody, in a suitable container.

### **Detailed Description**

The present invention relates to recombinant antigenic *S. pneumoniae* polypeptides and fragments thereof. The invention also relates to methods for using these polypeptides to produce immunological responses and to confer immunological protection to disease caused by members of the genus *Streptococcus*, at least isolates of the *S. pneumoniae* genus. The invention further relates to nucleic acid sequences which encode antigenic *S. pneumoniae* polypeptides and to methods for detecting *S. pneumoniae* nucleic acids and polypeptides in biological samples. The invention also relates to *S. pneumoniae*-specific antibodies and methods for detecting such antibodies produced in a host animal.

### **Definitions**

The following definitions are provided to clarify the subject matter which the inventors consider to be the present invention.

As used herein, the phrase "pathogenic agent" means an agent which causes a disease state or affliction in an animal. Included within this definition, for examples, are bacteria, protozoans, fungi, viruses and metazoan parasites which either produce a disease state or render an animal infected with such an organism susceptible to a disease state (*e.g.*, a secondary infection). Further included are species and strains of the genus *Streptococcus* which produce disease states in animals.

As used herein, the term "organism" means any living biological system, including viruses, regardless of whether it is a pathogenic agent.

As used herein, the term "*Streptococcus*" means any species or strain of bacteria which is members of the genus *Streptococcus*. Such species and strains are known to those of skill in the art, and include those that are pathogenic and those that are not.

As used herein, the phrase "one or more *S. pneumoniae* polypeptides of the present invention" means polypeptides comprising the amino acid sequence of one or more of the *S. pneumoniae* polypeptides described in Table 1 and disclosed as SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, and so on through SEQ ID NO:226. These polypeptides may be expressed as fusion proteins wherein the *S. pneumoniae* polypeptides of the present invention are linked to additional amino acid sequences which may be of streptococcal or non-streptococcal origin. This phrase further includes polypeptide comprising fragments of the *S. pneumoniae* polypeptides of the present invention.

Additional definitions are provided throughout the specification.

### ***Explanation of Table 1***

Table 1, below, provides information describing 113 open reading frames (ORFs) which encode potentially antigenic polypeptides of *S. pneumoniae* of the present invention. The table lists the ORF identifier which consists of the letters SP, which denote *S. pneumoniae*, followed immediately by a three digit numeric code, which arbitrarily number the potentially antigenic polypeptides of *S. pneumoniae* of the present invention and the nucleotide or amino acid sequence of each ORF and encoded polypeptide. The table further correlates the ORF identifier with a sequence identification number (SEQ ID NO:). The actual nucleotide or amino acid sequence of each ORF identifier is also shown in the Sequence Listing under the corresponding SEQ ID NO.

Thus, for example, the designation "SP126" refers to both the nucleotide and amino acid sequences of *S. pneumoniae* polypeptide number 126 of the present invention. Further, "SP126" correlates with the nucleotide

sequence shown as SEQ ID NO:223 and with the amino acid sequence shown as SEQ ID NO:224 as is described in Table 1.

The open reading frame within each "ORF" begins with the second nucleotide shown. Thus, the first codon for each nucleotide sequence shown is bases 2-4, the second 5-7, the third 8-10, and so on.

### ***Explanation of Table 2***

Table 2 lists the antigenic epitopes present in each of the *S. pneumoniae* polypeptides described in Table 1 as predicted by the inventors. Each *S. pneumoniae* polypeptide shown in Table 1 has one or more antigenic epitopes described in Table 2. It will be appreciated that depending on the analytical criteria used to predict antigenic determinants, the exact address of the determinant may vary slightly. The exact location of the antigenic determinant may shift by about 1 to 5 residues, more likely 1 to 2 residues, depending on the criteria used. Thus, the first antigenic determinant described in Table 2, "Lys-1 to Ile-10" of SP001, represents a peptide comprising the lysine at position 1 in SEQ ID NO:2 through and including the isoleucine at position 10 in SEQ ID NO:2, but may include more or fewer residues than those 10. It will also be appreciated that, generally speaking, amino acids can be added to either terminus of a peptide or polypeptide containing an antigenic epitope without affecting its activity, whereas removing residues from a peptide or polypeptide containing only the antigenic determinant is much more likely to destroy activity. It will be appreciated that the residues and locations shown described in Table 2 correspond to the amino acid sequences for each ORF shown in Table 1 and in the Sequence Listing.

### ***Explanation of Table 3***

Table 3 shows PCR primers designed by the inventors for the amplification of polynucleotides encoding polypeptides of the present invention according to the method of Example 1. PCR primer design is routine in the art and those shown in Table 3 are provided merely for the convenience of the skilled artisan. It will be appreciated that others can be used with equal success.

For each primer, the table lists the corresponding ORF designation from Table 1 followed by either an "A" or a "B". The "A" primers are the 5' primers and the "B" primers 3'. A restriction enzyme site was built into each primer to allow ease of cloning. The restriction enzyme which will recognize and cleave a sequence within each primer is shown in Table 3, as well, under the heading

"RE" for restriction enzyme. Finally the sequence identifier is shown in Table 3 for each primer for easy correlation with the Sequence Listing.

### *Selection of Nucleic Acid Sequences Encoding Antigenic S. pneumoniae Polypeptides*

The present invention provides a select number of ORFs from those presented in the fragments of the *S. pneumoniae* genome which may prove useful for the generation of a protective immune response. The sequenced *S. pneumoniae* genomic DNA was obtained from a sub-cultured isolate of *S. pneumoniae* Strain 7/87 14.8.91, which has been deposited at the American Type Culture Collection, as a convenience to those of skill in the art. The *S. pneumoniae* isolate was deposited on October 10, 1996 at the ATCC, 12301 Park Lawn Drive, Rockville, Maryland 20852, and given accession number 55840. A genomic library constructed from DNA isolated from the *S. pneumoniae* isolate was also deposited at the ATCC on October 11, 1996 and given ATCC Deposit No. 97755. A more complete listing of the sequence obtained from the *S. pneumoniae* genome may be found in co-pending U.S. Provisional Application Serial No. 60/029,960, filed 10/31/96, incorporated herein by reference in its entirety. Some ORFs contained in the subset of fragments of the *S. pneumoniae* genome disclosed herein were derived through the use of a number of screening criteria detailed below.

The selected ORFs do not consist of complete ORFs. Although a polypeptide representing a complete ORF may be the closest approximation of a protein native to an organism, it is not always preferred to express a complete ORF in a heterologous system. It may be challenging to express and purify a highly hydrophobic protein by common laboratory methods. Thus, the polypeptide vaccine candidates described herein may have been modified slightly to simplify the production of recombinant protein. For example, nucleotide sequences which encode highly hydrophobic domains, such as those found at the amino terminal signal sequence, have been excluded from some constructs used for *in vitro* expression of the polypeptides. Furthermore, any highly hydrophobic amino acid sequences occurring at the carboxy terminus have also been excluded from the recombinant expression constructs. Thus, in one embodiment, a polypeptide which represents a truncated or modified ORF may be used as an antigen.

While numerous methods are known in the art for selecting potentially immunogenic polypeptides, many of the ORFs disclosed herein were selected

on the basis of screening all theoretical *S. pneumoniae* ORFs for several aspects of potential immunogenicity. One set of selection criteria are as follows:

5 1. *Type I signal sequence*: An amino terminal type I signal sequence generally directs a nascent protein across the plasma and outer membranes to the exterior of the bacterial cell. Experimental evidence obtained from studies with *Escherichia coli* suggests that the typical type I signal sequence consists of the following biochemical and physical attributes (Izard, J. W. and Kendall, D. A. *Mol. Microbiol.* **13**:765-773 (1994)). The length of the type I signal sequence is approximately 15 to 25 primarily hydrophobic amino acid residues with a net positive charge in the extreme amino terminus. In addition, the central region of the signal sequence adopts an alpha-helical conformation in a hydrophobic environment. Finally, the region surrounding the actual site of cleavage is ideally six residues long, with small side-chain amino acids in the -1 and -3 positions.

15 2. *Type IV signal sequence*: The type IV signal sequence is an example of the several types of functional signal sequences which exist in addition to the type I signal sequence detailed above. Although functionally related, the type IV signal sequence possesses a unique set of biochemical and physical attributes (Strom, M. S. and Lory, S., *J. Bacteriol.* **174**:7345-7351 (1992)). These are typically six to eight amino acids with a net basic charge followed by an additional sixteen to thirty primarily hydrophobic residues. The cleavage site of a type IV signal sequence is typically after the initial six to eight amino acids at the extreme amino terminus. In addition, type IV signal sequences generally contain a phenylalanine residue at the +1 site relative to the cleavage site.

25 3. *Lipoprotein*: Studies of the cleavage sites of twenty-six bacterial lipoprotein precursors has allowed the definition of a consensus amino acid sequence for lipoprotein cleavage. Nearly three-fourths of the bacterial lipoprotein precursors examined contained the sequence L-(A,S)-(G,A)-C at positions -3 to +1, relative to the point of cleavage (Hayashi, S. and Wu, H. C., *J. Bioenerg. Biomembr.* **22**:451-471 (1990)).

30 4. *LPXTG motif*: It has been experimentally determined that most anchored proteins found on the surface of gram-positive bacteria possess a highly conserved carboxy terminal sequence. More than fifty such proteins from organisms such as *S. pyogenes*, *S. mutans*, *E. faecalis*, *S. pneumoniae*, and others, have been identified based on their extracellular location and carboxy terminal amino acid sequence (Fischetti, V. A., *ASM News* **62**:405-410 (1996)). The conserved region consists of six charged amino acids at the extreme carboxy terminus coupled to 15-20 hydrophobic amino acids

presumed to function as a transmembrane domain. Immediately adjacent to the transmembrane domain is a six amino acid sequence conserved in nearly all proteins examined. The amino acid sequence of this region is L-P-X-T-G-X, where X is any amino acid.

5 An algorithm for selecting antigenic and immunogenic *S. pneumoniae* polypeptides including the foregoing criteria was developed. Use of the algorithm by the inventors to select immunologically useful *S. pneumoniae* polypeptides resulted in the selection of a number of the disclosed ORFs. Polypeptides comprising the polypeptides identified in this group may be  
10 produced by techniques standard in the art and as further described herein.

### ***Nucleic Acid Molecules***

15 The present invention provides isolated nucleic acid molecules comprising polynucleotides encoding the *S. pneumoniae* polypeptides having the amino acid sequences described in Table 1 and shown as SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, and so on through SEQ ID NO:226, which were determined by sequencing the genome of *S. pneumoniae* and selected as putative immunogens.

20 Unless otherwise indicated, all nucleotide sequences determined by sequencing a DNA molecule herein were determined using an automated DNA sequencer (such as the Model 373 from Applied Biosystems, Inc.), and all amino acid sequences of polypeptides encoded by DNA molecules determined herein were predicted by translation of DNA sequences determined as above. Therefore, as is known in the art for any DNA sequence determined by this  
25 automated approach, any nucleotide sequence determined herein may contain some errors. Nucleotide sequences determined by automation are typically at least about 90% identical, more typically at least about 95% to at least about 99.9% identical to the actual nucleotide sequence of the sequenced DNA molecule. The actual sequence can be more precisely determined by other  
30 approaches including manual DNA sequencing methods well known in the art. As is also known in the art, a single insertion or deletion in a determined nucleotide sequence compared to the actual sequence will cause a frame shift in translation of the nucleotide sequence such that the predicted amino acid sequence encoded by a determined nucleotide sequence will be completely  
35 different from the amino acid sequence actually encoded by the sequenced DNA molecule, beginning at the point of such an insertion or deletion.

Unless otherwise indicated, each "nucleotide sequence" set forth herein is presented as a sequence of deoxyribonucleotides (abbreviated A, G, C and

T). However, by "nucleotide sequence" of a nucleic acid molecule or polynucleotide is intended, for a DNA molecule or polynucleotide, a sequence of deoxyribonucleotides, and for an RNA molecule or polynucleotide, the corresponding sequence of ribonucleotides (A, G, C and U), where each thymidine deoxyribonucleotide (T) in the specified deoxyribonucleotide sequence is replaced by the ribonucleotide uridine (U). For instance, reference to an RNA molecule having a sequence described in Table 1 set forth using deoxyribonucleotide abbreviations is intended to indicate an RNA molecule having a sequence in which each deoxyribonucleotide A, G or C described in Table 1 has been replaced by the corresponding ribonucleotide A, G or C, and each deoxyribonucleotide T has been replaced by a ribonucleotide U.

Nucleic acid molecules of the present invention may be in the form of RNA, such as mRNA, or in the form of DNA, including, for instance, cDNA and genomic DNA obtained by cloning or produced synthetically. The DNA may be double-stranded or single-stranded. Single-stranded DNA or RNA may be the coding strand, also known as the sense strand, or it may be the non-coding strand, also referred to as the anti-sense strand.

By "isolated" nucleic acid molecule(s) is intended a nucleic acid molecule, DNA or RNA, which has been removed from its native environment. For example, recombinant DNA molecules contained in a vector are considered isolated for the purposes of the present invention. Further examples of isolated DNA molecules include recombinant DNA molecules maintained in heterologous host cells or purified (partially or substantially) DNA molecules in solution. Isolated RNA molecules include *in vivo* or *in vitro* RNA transcripts of the DNA molecules of the present invention. Isolated nucleic acid molecules according to the present invention further include such molecules produced synthetically.

Isolated nucleic acid molecules of the present invention include DNA molecules comprising a nucleotide sequence described in Table 1 and shown as SEQ ID NO:1, SEQ ID NO:3, SEQ ID NO:5, and so on through SEQ ID NO:225; DNA molecules comprising the coding sequences for the polypeptides described in Table 1 and shown as SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, and so on through SEQ ID NO:226; and DNA molecules which comprise sequences substantially different from those described above but which, due to the degeneracy of the genetic code, still encode the *S. pneumoniae* polypeptides described in Table 1. Of course, the genetic code is well known in the art. Thus, it would be routine for one skilled in the art to generate such degenerate variants.



The invention also provides nucleic acid molecules having sequences complementary to any one of those described in Table 1. Such isolated molecules, particularly DNA molecules, are useful as probes for detecting expression of *Streptococcal* genes, for instance, by Northern blot analysis or the polymerase chain reaction (PCR).

The present invention is further directed to fragments of the isolated nucleic acid molecules described herein. By a fragment of an isolated nucleic acid molecule having a nucleotide sequence described in Table 1, is intended fragments at least about 15 nt, and more preferably at least about 17 nt, still more preferably at least about 20 nt, and even more preferably, at least about 25 nt in length which are useful as diagnostic probes and primers as discussed herein. Of course, larger fragments 50-100 nt in length are also useful according to the present invention as are fragments corresponding to most, if not all, of a nucleotide sequence described in Table 1. By a fragment at least 20 nt in length, for example, is intended fragments which include 20 or more contiguous bases of a nucleotide sequence as described in Table 1. Since the nucleotide sequences identified in Table 1 are provided as SEQ ID NO:1, SEQ ID NO:3, SEQ ID NO:5, and so on through SEQ ID NO:225, generating such DNA fragments would be routine to the skilled artisan. For example, such fragments could be generated synthetically.

Preferred nucleic acid fragments of the present invention also include nucleic acid molecules comprising nucleotide sequences encoding epitope-bearing portions of the *S. pneumoniae* polypeptides identified in Table 1. Such nucleic acid fragments of the present invention include, for example, nucleotide sequences encoding polypeptide fragments comprising from about the amino terminal residue to about the carboxy terminal residue of each fragment shown in Table 2. The above referred to polypeptide fragments are antigenic regions of the *S. pneumoniae* polypeptides identified in Table 1.

In another aspect, the invention provides isolated nucleic acid molecules comprising polynucleotides which hybridize under stringent hybridization conditions to a portion of a polynucleotide in a nucleic acid molecule of the invention described above, for instance, a nucleic acid sequence identified in Table 1. By "stringent hybridization conditions" is intended overnight incubation at 42°C in a solution comprising: 50% formamide, 5x SSC (150 mM NaCl, 15 mM trisodium citrate), 50 mM sodium phosphate (pH 7.6), 5x Denhardt's solution, 10% dextran sulfate, and 20 g/ml denatured, sheared salmon sperm DNA, followed by washing the filters in 0.1x SSC at about 65°C.

By polynucleotides which hybridize to a "portion" of a polynucleotide is intended polynucleotides (either DNA or RNA) which hybridize to at least about 15 nucleotides (nt), and more preferably at least about 17 nt, still more preferably at least about 20 nt, and even more preferably about 25-70 nt of the reference polynucleotide. These are useful as diagnostic probes and primers as discussed above and in more detail below.

Of course, polynucleotides hybridizing to a larger portion of the reference polynucleotide, for instance, a portion 50-100 nt in length, or even to the entire length of the reference polynucleotide, are also useful as probes according to the present invention, as are polynucleotides corresponding to most, if not all, of a nucleotide sequence as identified in Table 1. By a portion of a polynucleotide of "at least 20 nt in length," for example, is intended 20 or more contiguous nucleotides from the nucleotide sequence of the reference polynucleotide (e.g., a nucleotide sequences as described in Table 1). As noted above, such portions are useful diagnostically either as probes according to conventional DNA hybridization techniques or as primers for amplification of a target sequence by PCR, as described in the literature (for instance, in *Molecular Cloning, A Laboratory Manual*, 2nd. edition, Sambrook, J., Fritsch, E. F. and Maniatis, T., eds., Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y. (1989), the entire disclosure of which is hereby incorporated herein by reference).

Since nucleic acid sequences encoding the *S. pneumoniae* polypeptides of the present invention are identified in Table 1 and provided as SEQ ID NO:1, SEQ ID NO:3, SEQ ID NO:5, and so on through SEQ ID NO:225, generating polynucleotides which hybridize to portions of these sequences would be routine to the skilled artisan. For example, the hybridizing polynucleotides of the present invention could be generated synthetically according to known techniques.

As indicated, nucleic acid molecules of the present invention which encode *S. pneumoniae* polypeptides of the present invention may include, but are not limited to those encoding the amino acid sequences of the polypeptides by themselves; and additional coding sequences which code for additional amino acids, such as those which provide additional functionalities. Thus, the sequences encoding these polypeptides may be fused to a marker sequence, such as a sequence encoding a peptide which facilitates purification of the fused polypeptide. In certain preferred embodiments of this aspect of the invention, the marker amino acid sequence is a hexa-histidine peptide, such as the tag provided in a pQE vector (Qiagen, Inc.), among others, many of which are

commercially available. As described by Gentz and colleagues (*Proc. Natl. Acad. Sci. USA* **86**:821-824 (1989)), for instance, hexa-histidine provides for convenient purification of the resulting fusion protein.

Thus, the present invention also includes genetic fusions wherein the *S. pneumoniae* nucleic acid sequences coding sequences identified in Table 1 are linked to additional nucleic acid sequences to produce fusion proteins. These fusion proteins may include epitopes of streptococcal or non-streptococcal origin designed to produce proteins having enhanced immunogenicity. Further, the fusion proteins of the present invention may contain antigenic determinants known to provide helper T-cell stimulation, peptides encoding sites for post-translational modifications which enhance immunogenicity (*e.g.*, acylation), peptides which facilitate purification (*e.g.*, histidine "tag"), or amino acid sequences which target the fusion protein to a desired location (*e.g.*, a heterologous leader sequence).

In all cases of bacterial expression, an N-terminal methionine residues is added. In many cases, however, the N-terminal methionine residues is cleaved off post-translationally. Thus, the invention includes polypeptides shown in Table 1 with, and without an N-terminal methionine.

The present invention thus includes nucleic acid molecules and sequences which encode fusion proteins comprising one or more *S. pneumoniae* polypeptides of the present invention fused to an amino acid sequence which allows for post-translational modification to enhance immunogenicity. This post-translational modification may occur either *in vitro* or when the fusion protein is expressed *in vivo* in a host cell. An example of such a modification is the introduction of an amino acid sequence which results in the attachment of a lipid moiety.

Thus, as indicated above, the present invention includes genetic fusions wherein a *S. pneumoniae* nucleic acid sequence identified in Table 1 is linked to a nucleotide sequence encoding another amino acid sequence. These other amino acid sequences may be of streptococcal origin (*e.g.*, another sequence selected from Table 1) or non-streptococcal origin.

The present invention further relates to variants of the nucleic acid molecules of the present invention, which encode portions, analogs or derivatives of the *S. pneumoniae* polypeptides described in Table 1. Variants may occur naturally, such as a natural allelic variant. By an "allelic variant" is intended one of several alternate forms of a gene occupying a given locus on a chromosome of an organism (*Genes II*, Lewin, B., ed., John Wiley & Sons,

New York (1985)). Non-naturally occurring variants may be produced using art-known mutagenesis techniques.

Such variants include those produced by nucleotide substitutions, deletions or additions. The substitutions, deletions or additions may involve one or more nucleotides. These variants may be altered in coding regions, non-coding regions, or both. Alterations in the coding regions may produce conservative or non-conservative amino acid substitutions, deletions or additions. Especially preferred among these are silent substitutions, additions and deletions, which do not alter the properties and activities of the *S. pneumoniae* polypeptides disclosed herein or portions thereof. Silent substitution are most likely to be made in non-epitopic regions. Guidance regarding those regions containing epitopes is provided herein, for example, in Table 2. Also especially preferred in this regard are conservative substitutions.

Further embodiments of the invention include isolated nucleic acid molecules comprising a polynucleotide having a nucleotide sequence at least 90% identical, and more preferably at least 95%, 96%, 97%, 98% or 99% identical to: (a) a nucleotide sequence encoding any of the amino acid sequences of the polypeptides identified in Table 1; and (b) a nucleotide sequence complementary to any of the nucleotide sequences in (a) above.

By a polynucleotide having a nucleotide sequence at least, for example, 95% "identical" to a reference nucleotide sequence encoding a *S. pneumoniae* polypeptide described in Table 1, is intended that the nucleotide sequence of the polynucleotide is identical to the reference sequence except that the polynucleotide sequence may include up to five point mutations per each 100 nucleotides of the reference nucleotide sequence encoding the subject *S. pneumoniae* polypeptide. In other words, to obtain a polynucleotide having a nucleotide sequence at least 95% identical to a reference nucleotide sequence, up to 5% of the nucleotides in the reference sequence may be deleted or substituted with another nucleotide, or a number of nucleotides up to 5% of the total nucleotides in the reference sequence may be inserted into the reference sequence. These mutations of the reference sequence may occur at the 5' or 3' terminal positions of the reference nucleotide sequence or anywhere between those terminal positions, interspersed either individually among nucleotides in the reference sequence or in one or more contiguous groups within the reference sequence.

Certain nucleotides within some of the nucleic acid sequences shown in Table 1 were ambiguous upon sequencing. Completely unknown sequences are shown as an "N". Other unresolved nucleotides are known to be either a

purine, shown as "R", or a pyrimidine, shown as "Y". Accordingly, when determining identity between two nucleotide sequences, identity is met where any nucleotide, including an "R", "Y" or "N", is found in a test sequence and at the corresponding position in the reference sequence (from Table 1). Likewise, an A, G or "R" in a test sequence is identical to an "R" in the reference sequence; and a T, C or "Y" in a test sequence is identical to a "Y" in the reference sequence.

As a practical matter, whether any particular nucleic acid molecule is at least 90%, 95%, 96%, 97%, 98% or 99% identical to, for instance, a nucleotide sequence described in Table 1 can be determined conventionally using known computer programs such as the Bestfit program (Wisconsin Sequence Analysis Package, Version 8 for Unix, Genetics Computer Group, University Research Park, 575 Science Drive, Madison, WI 53711). Bestfit uses the local homology algorithm of Smith and Waterman (*Advances in Applied Mathematics* 2:482-489 (1981)), to find the best segment of homology between two sequences. When using Bestfit or any other sequence alignment program to determine whether a particular sequence is, for instance, 95% identical to a reference sequence according to the present invention, the parameters are set, of course, such that the percentage of identity is calculated over the full length of the reference nucleotide sequence and that gaps in homology of up to 5% of the total number of nucleotides in the reference sequence are allowed.

The present application is directed to nucleic acid molecules at least 90%, 95%, 96%, 97%, 98% or 99% identical to a nucleic acid sequences described in Table 1. One of skill in the art would still know how to use the nucleic acid molecule, for instance, as a hybridization probe or a polymerase chain reaction (PCR) primer. Uses of the nucleic acid molecules of the present invention include, *inter alia*, (1) isolating *Streptococcal* genes or allelic variants thereof from either a genomic or cDNA library and (2) Northern Blot or PCR analysis for detecting *Streptococcal* mRNA expression.

Of course, due to the degeneracy of the genetic code, one of ordinary skill in the art will immediately recognize that a large number of nucleic acid molecules having a sequence at least 90%, 95%, 96%, 97%, 98%, or 99% identical to a nucleic acid sequence identified in Table 1 will encode the same polypeptide. In fact, since degenerate variants of these nucleotide sequences all encode the same polypeptide, this will be clear to the skilled artisan even without performing the above described comparison assay.

It will be further recognized in the art that, for such nucleic acid molecules that are not degenerate variants, a reasonable number will also encode

5 proteins having antigenic epitopes of the *S. pneumoniae* polypeptides of the present invention. This is because the skilled artisan is fully aware of amino acid substitutions that are either less likely or not likely to significantly effect the antigenicity of a polypeptide (e.g., replacement of an amino acid in a region which is not believed to form an antigenic epitope). For example, since  
10 antigenic epitopes have been identified which contain as few as six amino acids (see Harlow, *et al.*, *Antibodies: A Laboratory Manual*, 2nd Ed.; Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York (1988), page 76), in instances where a polypeptide has multiple antigenic epitopes the alteration of several amino acid residues would often not be expected to eliminate all of the antigenic epitopes of that polypeptide. This is especially so when the alterations are in regions believed to not constitute antigenic epitopes.

### 15 *Vectors and Host Cells*

The present invention also relates to vectors which include the isolated DNA molecules of the present invention, host cells which are genetically engineered with the recombinant vectors, and the production of *S. pneumoniae* polypeptides or fragments thereof by recombinant techniques.

20 Recombinant constructs may be introduced into host cells using well known techniques such as infection, transduction, transfection, transvection, electroporation and transformation. The vector may be, for example, a phage, plasmid, viral or retroviral vector. Retroviral vectors may be replication competent or replication defective. In the latter case, viral propagation generally will occur only in complementing host cells.

25 The polynucleotides may be joined to a vector containing a selectable marker for propagation in a host. Generally, a plasmid vector is introduced in a precipitate, such as a calcium phosphate precipitate, or in a complex with a charged lipid. If the vector is a virus, it may be packaged *in vitro* using an appropriate packaging cell line and then transduced into host cells.

30 Preferred are vectors comprising *cis*-acting control regions to the polynucleotide of interest. Appropriate *trans*-acting factors may be supplied by the host, supplied by a complementing vector or supplied by the vector itself upon introduction into the host.

35 In certain preferred embodiments in this regard, the vectors provide for specific expression, which may be inducible and/or cell type-specific. Particularly preferred among such vectors are those inducible by environmental factors that are easy to manipulate, such as temperature and nutrient additives.

Expression vectors useful in the present invention include chromosomal-, episomal- and virus-derived vectors, *e.g.*, vectors derived from bacterial plasmids, bacteriophage, yeast episomes, yeast chromosomal elements, viruses such as baculoviruses, papova viruses, vaccinia viruses, adenoviruses, fowl pox viruses, pseudorabies viruses and retroviruses, and vectors derived from combinations thereof, such as cosmids and phagemids.

The DNA insert should be operatively linked to an appropriate promoter, such as the phage lambda PL promoter, the *E. coli lac*, *trp* and *tac* promoters, the SV40 early and late promoters and promoters of retroviral LTRs, to name a few. Other suitable promoters will be known to the skilled artisan. The expression constructs will further contain sites for transcription initiation, termination and, in the transcribed region, a ribosome binding site for translation. The coding portion of the mature transcripts expressed by the constructs will preferably include a translation initiating site at the beginning and a termination codon (UAA, UGA or UAG) appropriately positioned at the end of the polypeptide to be translated.

As indicated, the expression vectors will preferably include at least one selectable marker. Such markers include dihydrofolate reductase or neomycin resistance for eukaryotic cell culture and tetracycline or ampicillin resistance genes for culturing in *E. coli* and other bacteria. Representative examples of appropriate hosts include, but are not limited to, bacterial cells, such as *E. coli*, *Streptomyces* and *Salmonella typhimurium* cells; fungal cells, such as yeast cells; insect cells such as *Drosophila* S2 and *Spodoptera* Sf9 cells; animal cells such as CHO, COS and Bowes melanoma cells; and plant cells. Appropriate culture mediums and conditions for the above-described host cells are known in the art.

Among vectors preferred for use in bacteria include pQE70, pQE60 and pQE-9, available from Qiagen; pBS vectors, Phagescript vectors, Bluescript vectors, pNH8A, pNH16a, pNH18A, pNH46A available from Stratagene; pET series of vectors available from Novagen; and ptrc99a, pKK223-3, pKK233-3, pDR540, pRIT5 available from Pharmacia. Among preferred eukaryotic vectors are pWLNEO, pSV2CAT, pOG44, pXT1 and pSG available from Stratagene; and pSVK3, pBPV, pMSG and pSVL available from Pharmacia. Other suitable vectors will be readily apparent to the skilled artisan.

Among known bacterial promoters suitable for use in the present invention include the *E. coli lacI* and *lacZ* promoters, the T3 and T7 promoters, the *gpt* promoter, the lambda PR and PL promoters and the *trp* promoter. Suitable eukaryotic promoters include the CMV immediate early promoter, the

HSV thymidine kinase promoter, the early and late SV40 promoters, the promoters of retroviral LTRs, such as those of the Rous sarcoma virus (RSV), and metallothionein promoters, such as the mouse metallothionein-I promoter.

Introduction of the construct into the host cell can be effected by calcium phosphate transfection, DEAE-dextran mediated transfection, cationic lipid-mediated transfection, electroporation, transduction, infection or other methods. Such methods are described in many standard laboratory manuals (for example, Davis, *et al.*, *Basic Methods In Molecular Biology* (1986)).

Transcription of DNA encoding the polypeptides of the present invention by higher eukaryotes may be increased by inserting an enhancer sequence into the vector. Enhancers are *cis*-acting elements of DNA, usually about from 10 to 300 bp that act to increase transcriptional activity of a promoter in a given host cell-type. Examples of enhancers include the SV40 enhancer, which is located on the late side of the replication origin at bp 100 to 270, the cytomegalovirus early promoter enhancer, the polyoma enhancer on the late side of the replication origin, and adenovirus enhancers.

For secretion of the translated polypeptide into the lumen of the endoplasmic reticulum, into the periplasmic space or into the extracellular environment, appropriate secretion signals may be incorporated into the expressed polypeptide. The signals may be endogenous to the polypeptide or they may be heterologous signals.

The polypeptide may be expressed in a modified form, such as a fusion protein, and may include not only secretion signals, but also additional heterologous functional regions. For instance, a region of additional amino acids, particularly charged amino acids, may be added to the N-terminus of the polypeptide to improve stability and persistence in the host cell, during purification, or during subsequent handling and storage. Also, peptide moieties may be added to the polypeptide to facilitate purification. Such regions may be removed prior to final preparation of the polypeptide. The addition of peptide moieties to polypeptides to engender secretion or excretion, to improve stability and to facilitate purification, among others, are familiar and routine techniques in the art. A preferred fusion protein comprises a heterologous region from immunoglobulin that is useful to solubilize proteins. For example, EP-A-O 464 533 (Canadian counterpart 2045869) discloses fusion proteins comprising various portions of constant region of immunoglobulin molecules together with another human protein or part thereof. In many cases, the Fc part in a fusion protein is thoroughly advantageous for use in therapy and diagnosis and thus results, for example, in improved pharmacokinetic properties (EP-A 0232 262).



On the other hand, for some uses it would be desirable to be able to delete the Fc part after the fusion protein has been expressed, detected and purified in the advantageous manner described. This is the case when Fc portion proves to be a hindrance to use in therapy and diagnosis, for example when the fusion protein is to be used as antigen for immunizations. In drug discovery, for example, human proteins, such as, hIL-5-receptor has been fused with Fc portions for the purpose of high-throughput screening assays to identify antagonists of hIL-5. See Bennett, D. *et al.*, *J. Molec. Recogn.* 8:52-58 (1995) and Johanson, K. *et al.*, *J. Biol. Chem.* 270 (16):9459-9471 (1995).

The *S. pneumoniae* polypeptides can be recovered and purified from recombinant cell cultures by well-known methods including ammonium sulfate or ethanol precipitation, acid extraction, anion or cation exchange chromatography, phosphocellulose chromatography, hydrophobic interaction chromatography, affinity chromatography, hydroxylapatite chromatography, lectin chromatography and high performance liquid chromatography ("HPLC") is employed for purification. Polypeptides of the present invention include naturally purified products, products of chemical synthetic procedures, and products produced by recombinant techniques from a prokaryotic or eukaryotic host, including, for example, bacterial, yeast, higher plant, insect and mammalian cells.

### ***Polypeptides and Fragments***

The invention further provides isolated polypeptides having the amino acid sequences described in Table 1, and shown as SEQ ID NO:2, SEQ ID NO:4, SEQ ID NO:6, and so on through SEQ ID NO:226, and peptides or polypeptides comprising portions of the above polypeptides. The terms "peptide" and "oligopeptide" are considered synonymous (as is commonly recognized) and each term can be used interchangeably as the context requires to indicate a chain of at least two amino acids coupled by peptidyl linkages. The word "polypeptide" is used herein for chains containing more than ten amino acid residues. All oligopeptide and polypeptide formulas or sequences herein are written from left to right and in the direction from amino terminus to carboxy terminus.

Some amino acid sequences of the *S. pneumoniae* polypeptides described in Table 1 can be varied without significantly effecting the antigenicity of the polypeptides. If such differences in sequence are contemplated, it should be remembered that there will be critical areas on the polypeptide which determine antigenicity. In general, it is possible to replace residues which do

not form part of an antigenic epitope without significantly effecting the antigenicity of a polypeptide. Guidance for such alterations is given in Table 2 wherein epitopes for each polypeptide is delineated.

The polypeptides of the present invention are preferably provided in an isolated form. By "isolated polypeptide" is intended a polypeptide removed from its native environment. Thus, a polypeptide produced and/or contained within a recombinant host cell is considered isolated for purposes of the present invention. Also intended as an "isolated polypeptide" is a polypeptide that has been purified, partially or substantially, from a recombinant host cell. For example, recombinantly produced versions of the *S. pneumoniae* polypeptides described in Table 1 can be substantially purified by the one-step method described by Smith and Johnson (*Gene* 67:31-40 (1988)).

The polypeptides of the present invention include: (a) an amino acid sequence of any of the polypeptides described in Table 1; and (b) an amino acid sequence of an epitope-bearing portion of any one of the polypeptides of (a); as well as polypeptides with at least 70% similarity, and more preferably at least 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% similarity to those described in (a) or (b) above, as well as polypeptides having an amino acid sequence at least 70% identical, more preferably at least 75% identical, and still more preferably 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% identical to those above.

By "% similarity" for two polypeptides is intended a similarity score produced by comparing the amino acid sequences of the two polypeptides using the Bestfit program (Wisconsin Sequence Analysis Package, Version 8 for Unix, Genetics Computer Group, University Research Park, 575 Science Drive, Madison, WI 53711) and the default settings for determining similarity. Bestfit uses the local homology algorithm of Smith and Waterman (*Advances in Applied Mathematics* 2:482-489 (1981)) to find the best segment of similarity between two sequences.

By a polypeptide having an amino acid sequence at least, for example, 95% "identical" to a reference amino acid sequence of a *S. pneumoniae* polypeptide is intended that the amino acid sequence of the polypeptide is identical to the reference sequence except that the polypeptide sequence may include up to five amino acid alterations per each 100 amino acids of the reference amino acid sequence. In other words, to obtain a polypeptide having an amino acid sequence at least 95% identical to a reference amino acid sequence, up to 5% of the amino acid residues in the reference sequence may be deleted or substituted with another amino acid, or a number of amino acids up to

5 5% of the total amino acid residues in the reference sequence may be inserted into the reference sequence. These alterations of the reference sequence may occur at the amino or carboxy terminal positions of the reference amino acid sequence or anywhere between those terminal positions, interspersed either individually among residues in the reference sequence or in one or more contiguous groups within the reference sequence.

10 The amino acid sequences shown in Table 1 may have on or more "X" residues. "X" represents unknown. Thus, for purposes of defining identity, if any amino acid is present at the same position in a reference amino acid sequence (shown in Table 1) where an X is shown, the two sequences are identical at that position.

15 As a practical matter, whether any particular polypeptide is at least 70%, 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% identical to, for instance, an amino acid sequence shown in Table 1, can be determined conventionally using known computer programs such the Bestfit program (Wisconsin Sequence Analysis Package, Version 8 for Unix, Genetics Computer Group, University Research Park, 575 Science Drive, Madison, WI 53711). When using Bestfit or any other sequence alignment program to determine whether a particular sequence is, for instance, 95% identical to a reference sequence according to the present invention, the parameters are set, of course, such that the percentage of identity is calculated over the full length of the reference amino acid sequence and that gaps in homology of up to 5% of the total number of amino acid residues in the reference sequence are allowed.

25 As described below, the polypeptides of the present invention can also be used to raise polyclonal and monoclonal antibodies, which are useful in assays for detecting *Streptococcal* protein expression.

30 In another aspect, the invention provides peptides and polypeptides comprising epitope-bearing portions of the *S. pneumoniae* polypeptides of the invention. These epitopes are immunogenic or antigenic epitopes of the polypeptides of the invention. An "immunogenic epitope" is defined as a part of a protein that elicits an antibody response when the whole protein or polypeptide is the immunogen. These immunogenic epitopes are believed to be confined to a few loci on the molecule. On the other hand, a region of a protein molecule to which an antibody can bind is defined as an "antigenic determinant" or "antigenic epitope." The number of immunogenic epitopes of a protein generally is less than the number of antigenic epitopes (Geysen, *et al.*, *Proc. Natl. Acad. Sci. USA* **81**:3998- 4002 (1983)). Predicted antigenic epitopes are shown in Table 2, below.

As to the selection of peptides or polypeptides bearing an antigenic epitope (*i.e.*, that contain a region of a protein molecule to which an antibody can bind), it is well known in that art that relatively short synthetic peptides that mimic part of a protein sequence are routinely capable of eliciting an antiserum that reacts with the partially mimicked protein (for instance, Sutcliffe, J., *et al.*, *Science* **219**:660-666 (1983)). Peptides capable of eliciting protein-reactive sera are frequently represented in the primary sequence of a protein, can be characterized by a set of simple chemical rules, and are confined neither to immunodominant regions of intact proteins (*i.e.*, immunogenic epitopes) nor to the amino or carboxyl terminals. Peptides that are extremely hydrophobic and those of six or fewer residues generally are ineffective at inducing antibodies that bind to the mimicked protein; longer, peptides, especially those containing proline residues, usually are effective (Sutcliffe, *et al.*, *supra*, p. 661). For instance, 18 of 20 peptides designed according to these guidelines, containing 8-39 residues covering 75% of the sequence of the influenza virus hemagglutinin HA1 polypeptide chain, induced antibodies that reacted with the HA1 protein or intact virus; and 12/12 peptides from the MuLV polymerase and 18/18 from the rabies glycoprotein induced antibodies that precipitated the respective proteins.

Antigenic epitope-bearing peptides and polypeptides of the invention are therefore useful to raise antibodies, including monoclonal antibodies, that bind specifically to a polypeptide of the invention. Thus, a high proportion of hybridomas obtained by fusion of spleen cells from donors immunized with an antigen epitope-bearing peptide generally secrete antibody reactive with the native protein (Sutcliffe, *et al.*, *supra*, p. 663). The antibodies raised by antigenic epitope-bearing peptides or polypeptides are useful to detect the mimicked protein, and antibodies to different peptides may be used for tracking the fate of various regions of a protein precursor which undergoes post-translational processing. The peptides and anti-peptide antibodies may be used in a variety of qualitative or quantitative assays for the mimicked protein, for instance in competition assays since it has been shown that even short peptides (*e.g.*, about 9 amino acids) can bind and displace the larger peptides in immunoprecipitation assays (for instance, Wilson, *et al.*, *Cell* **37**:767-778 (1984) p. 777). The anti-peptide antibodies of the invention also are useful for purification of the mimicked protein, for instance, by adsorption chromatography using methods well known in the art.

Antigenic epitope-bearing peptides and polypeptides of the invention designed according to the above guidelines preferably contain a sequence of at

least seven, more preferably at least nine and most preferably between about 15 to about 30 amino acids contained within the amino acid sequence of a polypeptide of the invention. However, peptides or polypeptides comprising a larger portion of an amino acid sequence of a polypeptide of the invention, containing about 30 to about 50 amino acids, or any length up to and including the entire amino acid sequence of a polypeptide of the invention, also are considered epitope-bearing peptides or polypeptides of the invention and also are useful for inducing antibodies that react with the mimicked protein. Preferably, the amino acid sequence of the epitope-bearing peptide is selected to provide substantial solubility in aqueous solvents (*i.e.*, the sequence includes relatively hydrophilic residues and highly hydrophobic sequences are preferably avoided); and sequences containing proline residues are particularly preferred.

Non-limiting examples of antigenic polypeptides or peptides that can be used to generate *Streptococcal*-specific antibodies include portions of the amino acid sequences identified in Table 1. More specifically, Table 2 discloses antigenic fragments of polypeptides of the present invention, which antigenic fragments comprise amino acid sequences from about the first amino acid residues indicated to about the last amino acid residue indicated for each fragment. The polypeptide fragments disclosed in Table 2 are believed to be antigenic regions of the *S. pneumoniae* polypeptides described in Table 1. Thus the invention further includes isolated peptides and polypeptides comprising an amino acid sequence of an epitope shown in Table 2 and polynucleotides encoding said polypeptides.

The epitope-bearing peptides and polypeptides of the invention may be produced by any conventional means for making peptides or polypeptides including recombinant means using nucleic acid molecules of the invention. For instance, an epitope-bearing amino acid sequence of the present invention may be fused to a larger polypeptide which acts as a carrier during recombinant production and purification, as well as during immunization to produce anti-peptide antibodies. Epitope-bearing peptides also may be synthesized using known methods of chemical synthesis. For instance, Houghten has described a simple method for synthesis of large numbers of peptides, such as 10-20 mg of 248 different 13 residue peptides representing single amino acid variants of a segment of the HA1 polypeptide which were prepared and characterized (by ELISA-type binding studies) in less than four weeks (Houghten, R. A. Proc. Natl. Acad. Sci. USA **82**:5131-5135 (1985)). This "Simultaneous Multiple Peptide Synthesis (SMPS)" process is further described in U.S. Patent No. 4,631,211 to Houghten and coworkers (1986). In this procedure the individual

resins for the solid-phase synthesis of various peptides are contained in separate solvent-permeable packets, enabling the optimal use of the many identical repetitive steps involved in solid-phase methods. A completely manual procedure allows 500-1000 or more syntheses to be conducted simultaneously (Houghten, *et al.*, *supra*, p. 5134).

Epitope-bearing peptides and polypeptides of the invention are used to induce antibodies according to methods well known in the art (for instance, Sutcliffe, *et al.*, *supra*; Wilson, *et al.*, *supra*; Chow, M., *et al.*, *Proc. Natl. Acad. Sci. USA* **82**:910-914; and Bittle, F. J., *et al.*, *J. Gen. Virol.* **66**:2347-2354 (1985)). Generally, animals may be immunized with free peptide; however, anti-peptide antibody titer may be boosted by coupling of the peptide to a macromolecular carrier, such as keyhole limpet hemacyanin (KLH) or tetanus toxoid. For instance, peptides containing cysteine may be coupled to carrier using a linker such as m-maleimidobenzoyl-N-hydroxysuccinimide ester (MBS), while other peptides may be coupled to carrier using a more general linking agent such as glutaraldehyde. Animals such as rabbits, rats and mice are immunized with either free or carrier-coupled peptides, for instance, by intraperitoneal and/or intradermal injection of emulsions containing about 100  $\mu$ g peptide or carrier protein and Freund's adjuvant. Several booster injections may be needed, for instance, at intervals of about two weeks, to provide a useful titer of anti-peptide antibody which can be detected, for example, by ELISA assay using free peptide adsorbed to a solid surface. The titer of anti-peptide antibodies in serum from an immunized animal may be increased by selection of anti-peptide antibodies, for instance, by adsorption to the peptide on a solid support and elution of the selected antibodies according to methods well known in the art.

Immunogenic epitope-bearing peptides of the invention, *i.e.*, those parts of a protein that elicit an antibody response when the whole protein is the immunogen, are identified according to methods known in the art. For instance, Geysen, *et al.*, *supra*, discloses a procedure for rapid concurrent synthesis on solid supports of hundreds of peptides of sufficient purity to react in an enzyme-linked immunosorbent assay. Interaction of synthesized peptides with antibodies is then easily detected without removing them from the support. In this manner a peptide bearing an immunogenic epitope of a desired protein may be identified routinely by one of ordinary skill in the art. For instance, the immunologically important epitope in the coat protein of foot-and-mouth disease virus was located by Geysen *et al. supra* with a resolution of seven amino acids by synthesis of an overlapping set of all 208 possible hexapeptides covering the

entire 213 amino acid sequence of the protein. Then, a complete replacement set of peptides in which all 20 amino acids were substituted in turn at every position within the epitope were synthesized, and the particular amino acids conferring specificity for the reaction with antibody were determined. Thus, peptide analogs of the epitope-bearing peptides of the invention can be made routinely by this method. U.S. Patent No. 4,708,781 to Geysen (1987) further describes this method of identifying a peptide bearing an immunogenic epitope of a desired protein.

Further still, U.S. Patent No. 5,194,392, to Geysen (1990), describes a general method of detecting or determining the sequence of monomers (amino acids or other compounds) which is a topological equivalent of the epitope (*i.e.*, a "mimotope") which is complementary to a particular paratope (antigen binding site) of an antibody of interest. More generally, U.S. Patent No. 4,433,092, also to Geysen (1989), describes a method of detecting or determining a sequence of monomers which is a topographical equivalent of a ligand which is complementary to the ligand binding site of a particular receptor of interest. Similarly, U.S. Patent No. 5,480,971 to Houghten, R. A. *et al.* (1996) discloses linear C<sub>1</sub>-C<sub>7</sub>-alkyl peralkylated oligopeptides and sets and libraries of such peptides, as well as methods for using such oligopeptide sets and libraries for determining the sequence of a peralkylated oligopeptide that preferentially binds to an acceptor molecule of interest. Thus, non-peptide analogs of the epitope-bearing peptides of the invention also can be made routinely by these methods.

The entire disclosure of each document cited in this section on "Polypeptides and Fragments" is hereby incorporated herein by reference.

As one of skill in the art will appreciate, the polypeptides of the present invention and the epitope-bearing fragments thereof described above can be combined with parts of the constant domain of immunoglobulins (IgG), resulting in chimeric polypeptides. These fusion proteins facilitate purification and show an increased half-life *in vivo*. This has been shown, *e.g.*, for chimeric proteins consisting of the first two domains of the human CD4-polypeptide and various domains of the constant regions of the heavy or light chains of mammalian immunoglobulins (EPA 0,394,827; Traunecker *et al.*, *Nature* 331:84-86 (1988)). Fusion proteins that have a disulfide-linked dimeric structure due to the IgG part can also be more efficient in binding and neutralizing other molecules than a monomeric *S. pneumoniae* polypeptide or fragment thereof alone (Fountoulakis *et al.*, *J. Biochem.* 270:3958-3964 (1995)).

### Diagnostic Assays

The present invention further relates to a method for assaying for *Streptococcal* infection in an animal *via* detecting the expression of genes encoding *Streptococcal* polypeptides (*e.g.*, the polypeptides described Table 1). This method comprises analyzing tissue or body fluid from the animal for *Streptococcus*-specific antibodies or *Streptococcal* nucleic acids or proteins. Analysis of nucleic acid specific to *Streptococcus* can be done by PCR or hybridization techniques using nucleic acid sequences of the present invention as either hybridization probes or primers (*cf. Molecular Cloning: A Laboratory Manual, second edition*, edited by Sambrook, Fritsch, & Maniatis, Cold Spring Harbor Laboratory, 1989; Ereemeeva *et al.*, *J. Clin. Microbiol.* 32:803-810 (1994) which describes differentiation among spotted fever group *Rickettsiae* species by analysis of restriction fragment length polymorphism of PCR-amplified DNA). Methods for detecting *B. burgdorferi* nucleic acids *via* PCR are described, for example, in Chen *et al.*, *J. Clin. Microbiol.* 32:589-595 (1994).

Where diagnosis of a disease state related to infection with *Streptococcus* has already been made, the present invention is useful for monitoring progression or regression of the disease state whereby patients exhibiting enhanced *Streptococcus* gene expression will experience a worse clinical outcome relative to patients expressing these gene(s) at a lower level.

By "assaying for *Streptococcal* infection in an animal *via* detection of genes encoding *Streptococcal* polypeptides" is intended qualitatively or quantitatively measuring or estimating the level of one or more *Streptococcus* polypeptides or the level of nucleic acid encoding *Streptococcus* polypeptides in a first biological sample either directly (*e.g.*, by determining or estimating absolute protein level or nucleic level) or relatively (*e.g.*, by comparing to the *Streptococcus* polypeptide level or mRNA level in a second biological sample). The *Streptococcus* polypeptide level or nucleic acid level in the second sample used for a relative comparison may be undetectable if obtained from an animal which is not infected with *Streptococcus*. When monitoring the progression or regression of a disease state, the *Streptococcus* polypeptide level or nucleic acid level may be compared to a second sample obtained from either an animal infected with *Streptococcus* or the same animal from which the first sample was obtained but taken from that animal at a different time than the first. As will be appreciated in the art, once a standard *Streptococcus* polypeptide level or nucleic



acid level which corresponds to a particular stage of a *Streptococcus* infection is known, it can be used repeatedly as a standard for comparison.

By "biological sample" is intended any biological sample obtained from an animal, cell line, tissue culture, or other source which contains *Streptococcus* polypeptide, mRNA, or DNA. Biological samples include body fluids (such as plasma and synovial fluid) which contain *Streptococcus* polypeptides, and muscle, skin, and cartilage tissues. Methods for obtaining tissue biopsies and body fluids are well known in the art.

The present invention is useful for detecting diseases related to *Streptococcus* infections in animals. Preferred animals include monkeys, apes, cats, dogs, cows, pigs, mice, horses, rabbits and humans. Particularly preferred are humans.

Total RNA can be isolated from a biological sample using any suitable technique such as the single-step guanidinium-thiocyanate-phenol-chloroform method described in Chomczynski and Sacchi, *Anal. Biochem.* 162:156-159 (1987). mRNA encoding *Streptococcus* polypeptides having sufficient homology to the nucleic acid sequences identified in Table 1 to allow for hybridization between complementary sequences are then assayed using any appropriate method. These include Northern blot analysis, S1 nuclease mapping, the polymerase chain reaction (PCR), reverse transcription in combination with the polymerase chain reaction (RT-PCR), and reverse transcription in combination with the ligase chain reaction (RT-LCR).

Northern blot analysis can be performed as described in Harada *et al.*, *Cell* 63:303-312 (1990). Briefly, total RNA is prepared from a biological sample as described above. For the Northern blot, the RNA is denatured in an appropriate buffer (such as glyoxal/dimethyl sulfoxide/sodium phosphate buffer), subjected to agarose gel electrophoresis, and transferred onto a nitrocellulose filter. After the RNAs have been linked to the filter by a UV linker, the filter is prehybridized in a solution containing formamide, SSC, Denhardt's solution, denatured salmon sperm, SDS, and sodium phosphate buffer. A *S. pneumoniae* polypeptide DNA sequence shown in Table 1 labeled according to any appropriate method (such as the <sup>32</sup>P-multiprimered DNA labeling system (Amersham)) is used as probe. After hybridization overnight, the filter is washed and exposed to x-ray film. DNA for use as probe according to the present invention is described in the sections above and will preferably at least 15 bp in length.

S1 mapping can be performed as described in Fujita *et al.*, *Cell* 49:357-367 (1987). To prepare probe DNA for use in S1 mapping, the sense

strand of an above-described *S. pneumoniae* DNA sequence of the present invention is used as a template to synthesize labeled antisense DNA. The antisense DNA can then be digested using an appropriate restriction endonuclease to generate further DNA probes of a desired length. Such antisense probes are useful for visualizing protected bands corresponding to the target mRNA (*i.e.*, mRNA encoding *Streptococcus* polypeptides).

Preferably, levels of mRNA encoding *Streptococcus* polypeptides are assayed using the RT-PCR method described in Makino *et al.*, *Technique* 2:295-301 (1990). By this method, the radioactivities of the "amplicons" in the polyacrylamide gel bands are linearly related to the initial concentration of the target mRNA. Briefly, this method involves adding total RNA isolated from a biological sample in a reaction mixture containing a RT primer and appropriate buffer. After incubating for primer annealing, the mixture can be supplemented with a RT buffer, dNTPs, DTT, RNase inhibitor and reverse transcriptase. After incubation to achieve reverse transcription of the RNA, the RT products are then subject to PCR using labeled primers. Alternatively, rather than labeling the primers, a labeled dNTP can be included in the PCR reaction mixture. PCR amplification can be performed in a DNA thermal cycler according to conventional techniques. After a suitable number of rounds to achieve amplification, the PCR reaction mixture is electrophoresed on a polyacrylamide gel. After drying the gel, the radioactivity of the appropriate bands (corresponding to the mRNA encoding the *Streptococcus* polypeptides)) is quantified using an imaging analyzer. RT and PCR reaction ingredients and conditions, reagent and gel concentrations, and labeling methods are well known in the art. Variations on the RT-PCR method will be apparent to the skilled artisan.

Assaying *Streptococcus* polypeptide levels in a biological sample can occur using any art-known method. Preferred for assaying *Streptococcus* polypeptide levels in a biological sample are antibody-based techniques. For example, *Streptococcus* polypeptide expression in tissues can be studied with classical immunohistological methods. In these, the specific recognition is provided by the primary antibody (polyclonal or monoclonal) but the secondary detection system can utilize fluorescent, enzyme, or other conjugated secondary antibodies. As a result, an immunohistological staining of tissue section for pathological examination is obtained. Tissues can also be extracted, *e.g.*, with urea and neutral detergent, for the liberation of *Streptococcus* polypeptides for Western-blot or dot/slot assay (Jalkanen, M., *et al.*, *J. Cell. Biol.* 101:976-985 (1985); Jalkanen, M., *et al.*, *J. Cell. Biol.* 105:3087-3096 (1987)). In this

technique, which is based on the use of cationic solid phases, quantitation of a *Streptococcus* polypeptide can be accomplished using an isolated *Streptococcus* polypeptide as a standard. This technique can also be applied to body fluids.

Other antibody-based methods useful for detecting *Streptococcus* polypeptide gene expression include immunoassays, such as the enzyme linked immunosorbent assay (ELISA) and the radioimmunoassay (RIA). For example, a *Streptococcus* polypeptide-specific monoclonal antibodies can be used both as an immunoabsorbent and as an enzyme-labeled probe to detect and quantify a *Streptococcus* polypeptide. The amount of a *Streptococcus* polypeptide present in the sample can be calculated by reference to the amount present in a standard preparation using a linear regression computer algorithm. Such an ELISA for detecting a tumor antigen is described in Iacobelli *et al.*, *Breast Cancer Research and Treatment* 11:19-30 (1988). In another ELISA assay, two distinct specific monoclonal antibodies can be used to detect *Streptococcus* polypeptides in a body fluid. In this assay, one of the antibodies is used as the immunoabsorbent and the other as the enzyme-labeled probe.

The above techniques may be conducted essentially as a "one-step" or "two-step" assay. The "one-step" assay involves contacting the *Streptococcus* polypeptide with immobilized antibody and, without washing, contacting the mixture with the labeled antibody. The "two-step" assay involves washing before contacting the mixture with the labeled antibody. Other conventional methods may also be employed as suitable. It is usually desirable to immobilize one component of the assay system on a support, thereby allowing other components of the system to be brought into contact with the component and readily removed from the sample.

*Streptococcus* polypeptide-specific antibodies for use in the present invention can be raised against an intact *S. pneumoize* polypeptide of the present invention or fragment thereof. These polypeptides and fragments may be administered to an animal (*e.g.*, rabbit or mouse) either with a carrier protein (*e.g.*, albumin) or, if long enough (*e.g.*, at least about 25 amino acids), without a carrier.

As used herein, the term "antibody" (Ab) or "monoclonal antibody" (Mab) is meant to include intact molecules as well as antibody fragments (such as, for example, Fab and  $F(ab')_2$  fragments) which are capable of specifically binding to a *Streptococcus* polypeptide. Fab and  $F(ab')_2$  fragments lack the Fc fragment of intact antibody, clear more rapidly from the circulation, and may have less non-specific tissue binding of an intact antibody (Wahl *et al.*, *J. Nucl. Med.* 24:316-325 (1983)). Thus, these fragments are preferred.

The antibodies of the present invention may be prepared by any of a variety of methods. For example, the *S. pneumoniae* polypeptides identified in Table 1, or fragments thereof, can be administered to an animal in order to induce the production of sera containing polyclonal antibodies. In a preferred method, a preparation of a *S. pneumoniae* polypeptide of the present invention is prepared and purified to render it substantially free of natural contaminants. Such a preparation is then introduced into an animal in order to produce polyclonal antisera of high specific activity.

In the most preferred method, the antibodies of the present invention are monoclonal antibodies. Such monoclonal antibodies can be prepared using hybridoma technology (Kohler *et al.*, *Nature* 256:495 (1975); Kohler *et al.*, *Eur. J. Immunol.* 6:511 (1976); Kohler *et al.*, *Eur. J. Immunol.* 6:292 (1976); Hammerling *et al.*, In: *Monoclonal Antibodies and T-Cell Hybridomas*, Elsevier, N.Y., (1981) pp. 563-681 ). In general, such procedures involve immunizing an animal (preferably a mouse) with a *S. pneumoniae* polypeptide antigen of the present invention. Suitable cells can be recognized by their capacity to bind anti-*Streptococcus* polypeptide antibody. Such cells may be cultured in any suitable tissue culture medium; however, it is preferable to culture cells in Earle's modified Eagle's medium supplemented with 10% fetal bovine serum (inactivated at about 56°C), and supplemented with about 10 g/l of nonessential amino acids, about 1,000 U/ml of penicillin, and about 100 µg/ml of streptomycin. The splenocytes of such mice are extracted and fused with a suitable myeloma cell line. Any suitable myeloma cell line may be employed in accordance with the present invention; however, it is preferable to employ the parent myeloma cell line (SP<sub>2</sub>O), available from the American Type Culture Collection, Rockville, Maryland. After fusion, the resulting hybridoma cells are selectively maintained in HAT medium, and then cloned by limiting dilution as described by Wands *et al.* (*Gastroenterology* 80:225-232 (1981)). The hybridoma cells obtained through such a selection are then assayed to identify clones which secrete antibodies capable of binding the *Streptococcus* polypeptide antigen administered to immunized animal.

Alternatively, additional antibodies capable of binding to *Streptococcus* polypeptide antigens may be produced in a two-step procedure through the use of anti-idiotypic antibodies. Such a method makes use of the fact that antibodies are themselves antigens, and that, therefore, it is possible to obtain an antibody which binds to a second antibody. In accordance with this method, *Streptococcus* polypeptide-specific antibodies are used to immunize an animal, preferably a mouse. The splenocytes of such an animal are then used to

produce hybridoma cells, and the hybridoma cells are screened to identify clones which produce an antibody whose ability to bind to the *Streptococcus* polypeptide-specific antibody can be blocked by a *Streptococcus* polypeptide antigen. Such antibodies comprise anti-idiotypic antibodies to the *Streptococcus* polypeptide-specific antibody and can be used to immunize an animal to induce formation of further *Streptococcus* polypeptide-specific antibodies.

It will be appreciated that Fab and F(ab')<sub>2</sub> and other fragments of the antibodies of the present invention may be used according to the methods disclosed herein. Such fragments are typically produced by proteolytic cleavage, using enzymes such as papain (to produce Fab fragments) or pepsin (to produce F(ab')<sub>2</sub> fragments). Alternatively, *Streptococcus* polypeptide-binding fragments can be produced through the application of recombinant DNA technology or through synthetic chemistry.

Of special interest to the present invention are antibodies to *Streptococcus* polypeptide antigens which are produced in humans, or are "humanized" (i.e., non-immunogenic in a human) by recombinant or other technology. Humanized antibodies may be produced, for example by replacing an immunogenic portion of an antibody with a corresponding, but non-immunogenic portion (i.e., chimeric antibodies) (Robinson, R.R. *et al.*, International Patent Publication PCT/US86/02269; Akira, K. *et al.*, European Patent Application 184,187; Taniguchi, M., European Patent Application 171,496; Morrison, S.L. *et al.*, European Patent Application 173,494; Neuberger, M.S. *et al.*, PCT Application WO 86/01533; Cabilly, S. *et al.*, European Patent Application 125,023; Better, M. *et al.*, *Science* 240:1041-1043 (1988); Liu, A.Y. *et al.*, *Proc. Natl. Acad. Sci. USA* 84:3439-3443 (1987); Liu, A.Y. *et al.*, *J. Immunol.* 139:3521-3526 (1987); Sun, L.K. *et al.*, *Proc. Natl. Acad. Sci. USA* 84:214-218 (1987); Nishimura, Y. *et al.*, *Canc. Res.* 47:999-1005 (1987); Wood, C.R. *et al.*, *Nature* 314:446-449 (1985); Shaw *et al.*, *J. Natl. Cancer Inst.* 80:1553-1559 (1988). General reviews of "humanized" chimeric antibodies are provided by Morrison, S.L. (*Science*, 229:1202-1207 (1985)) and by Oi, V.T. *et al.*, *BioTechniques* 4:214 (1986)). Suitable "humanized" antibodies can be alternatively produced by CDR or CEA substitution (Jones, P.T. *et al.*, *Nature* 321:552-525 (1986); Verhoeyan *et al.*, *Science* 239:1534 (1988); Beidler, C.B. *et al.*, *J. Immunol.* 141:4053-4060 (1988)).

Suitable enzyme labels include, for example, those from the oxidase group, which catalyze the production of hydrogen peroxide by reacting with substrate. Glucose oxidase is particularly preferred as it has good stability and

its substrate (glucose) is readily available. Activity of an oxidase label may be assayed by measuring the concentration of hydrogen peroxide formed by the enzyme-labeled antibody/substrate reaction. Besides enzymes, other suitable labels include radioisotopes, such as iodine ( $^{125}\text{I}$ ,  $^{121}\text{I}$ ), carbon ( $^{14}\text{C}$ ), sulphur ( $^{35}\text{S}$ ), tritium ( $^3\text{H}$ ), indium ( $^{112}\text{In}$ ), and technetium ( $^{99\text{m}}\text{Tc}$ ), and fluorescent labels, such as fluorescein and rhodamine, and biotin.

Further suitable labels for the *Streptococcus* polypeptide-specific antibodies of the present invention are provided below. Examples of suitable enzyme labels include malate dehydrogenase, staphylococcal nuclease, delta-5-steroid isomerase, yeast-alcohol dehydrogenase, alpha-glycerol phosphate dehydrogenase, triose phosphate isomerase, peroxidase, alkaline phosphatase, asparaginase, glucose oxidase, beta-galactosidase, ribonuclease, urease, catalase, glucose-6-phosphate dehydrogenase, glucoamylase, and acetylcholine esterase.

Examples of suitable radioisotopic labels include  $^3\text{H}$ ,  $^{111}\text{In}$ ,  $^{125}\text{I}$ ,  $^{131}\text{I}$ ,  $^{32}\text{P}$ ,  $^{35}\text{S}$ ,  $^{14}\text{C}$ ,  $^{51}\text{Cr}$ ,  $^{57}\text{Co}$ ,  $^{58}\text{Co}$ ,  $^{59}\text{Fe}$ ,  $^{75}\text{Se}$ ,  $^{152}\text{Eu}$ ,  $^{90}\text{Y}$ ,  $^{67}\text{Cu}$ ,  $^{217}\text{Bi}$ ,  $^{211}\text{At}$ ,  $^{212}\text{Pb}$ ,  $^{47}\text{Sc}$ ,  $^{109}\text{Pd}$ , etc.  $^{111}\text{In}$  is a preferred isotope where *in vivo* imaging is used since it avoids the problem of dehalogenation of the  $^{125}\text{I}$  or  $^{131}\text{I}$ -labeled monoclonal antibody by the liver. In addition, this radionucleotide has a more favorable gamma emission energy for imaging (Perkins *et al.*, *Eur. J. Nucl. Med.* 10:296-301 (1985); Carasquillo *et al.*, *J. Nucl. Med.* 28:281-287 (1987)). For example,  $^{111}\text{In}$  coupled to monoclonal antibodies with 1-(P-isothiocyanatobenzyl)-DPTA has shown little uptake in non-tumorous tissues, particularly the liver, and therefore enhances specificity of tumor localization (Esteban *et al.*, *J. Nucl. Med.* 28:861-870 (1987)).

Examples of suitable non-radioactive isotopic labels include  $^{157}\text{Gd}$ ,  $^{55}\text{Mn}$ ,  $^{162}\text{Dy}$ ,  $^{52}\text{Tr}$ , and  $^{56}\text{Fe}$ .

Examples of suitable fluorescent labels include an  $^{152}\text{Eu}$  label, a fluorescein label, an isothiocyanate label, a rhodamine label, a phycoerythrin label, a phycocyanin label, an allophycocyanin label, an o-phthaldehyde label, and a fluorecamine label.

Examples of suitable toxin labels include diphtheria toxin, ricin, and cholera toxin.

Examples of chemiluminescent labels include a luminal label, an isoluminal label, an aromatic acridinium ester label, an imidazole label, an acridinium salt label, an oxalate ester label, a luciferin label, a luciferase label, and an aequorin label.

Examples of nuclear magnetic resonance contrasting agents include heavy metal nuclei such as Gd, Mn, and iron.

Typical techniques for binding the above-described labels to antibodies are provided by Kennedy *et al.*, *Clin. Chim. Acta* 70:1-31 (1976), and Schurs *et al.*, *Clin. Chim. Acta* 81:1-40 (1977). Coupling techniques mentioned in the latter are the glutaraldehyde method, the periodate method, the dimaleimide method, the m-maleimidobenzyl-N-hydroxy-succinimide ester method, all of which methods are incorporated by reference herein.

In a related aspect, the invention includes a diagnostic kit for use in screening serum containing antibodies specific against *S. pneumoniae* infection. Such a kit may include an isolated *S. pneumoniae* antigen comprising an epitope which is specifically immunoreactive with at least one anti-*S. pneumoniae* antibody. Such a kit also includes means for detecting the binding of said antibody to the antigen. In specific embodiments, the kit may include a recombinantly produced or chemically synthesized peptide or polypeptide antigen. The peptide or polypeptide antigen may be attached to a solid support.

In a more specific embodiment, the detecting means of the above-described kit includes a solid support to which said peptide or polypeptide antigen is attached. Such a kit may also include a non-attached reporter-labelled anti-human antibody. In this embodiment, binding of the antibody to the *S. pneumoniae* antigen can be detected by binding of the reporter labelled antibody to the anti-*S. pneumoniae* antibody.

In a related aspect, the invention includes a method of detecting *S. pneumoniae* infection in a subject. This detection method includes reacting a body fluid, preferably serum, from the subject with an isolated *S. pneumoniae* antigen, and examining the antigen for the presence of bound antibody. In a specific embodiment, the method includes a polypeptide antigen attached to a solid support, and serum is reacted with the support. Subsequently, the support is reacted with a reporter-labelled anti-human antibody. The support is then examined for the presence of reporter-labelled antibody.

The solid surface reagent employed in the above assays and kits is prepared by known techniques for attaching protein material to solid support material, such as polymeric beads, dip sticks, 96-well plates or filter material. These attachment methods generally include non-specific adsorption of the protein to the support or covalent attachment of the protein, typically through a free amine group, to a chemically reactive group on the solid support, such as an activated carboxyl, hydroxyl, or aldehyde group. Alternatively, streptavidin coated plates can be used in conjunction with biotinylated antigen(s).

### *Therapeutics and Modes of Administration*

5 The present invention also provides vaccines comprising one or more polypeptides of the present invention. Heterogeneity in the composition of a vaccine may be provided by combining *S. pneumoniae* polypeptides of the present invention. Multi-component vaccines of this type are desirable because they are likely to be more effective in eliciting protective immune responses against multiple species and strains of the *Streptococcus* genus than single polypeptide vaccines. Thus, as discussed in detail below, a multi-component vaccine of the present invention may contain one or more, preferably 2 to about 20, more preferably 2 to about 15, and most preferably 3 to about 8, of the *S. pneumoniae* polypeptides identified in Table 1, or fragments thereof.

10 Multi-component vaccines are known in the art to elicit antibody production to numerous immunogenic components. Decker, M. and Edwards, K., *J. Infect. Dis.* 174:S270-275 (1996). In addition, a hepatitis B, diphtheria, tetanus, pertussis tetravalent vaccine has recently been demonstrated to elicit protective levels of antibodies in human infants against all four pathogenic agents. Aristegui, J. *et al.*, *Vaccine* 15:7-9 (1997).

15 The present invention thus also includes multi-component vaccines. These vaccines comprise more than one polypeptide, immunogen or antigen. An example of such a multi-component vaccine would be a vaccine comprising more than one of the *S. pneumoniae* polypeptides described in Table 1. A second example is a vaccine comprising one or more, for example 2 to 10, of the *S. pneumoniae* polypeptides identified in Table 1 and one or more, for example 2 to 10, additional polypeptides of either streptococcal or non-streptococcal origin. Thus, a multi-component vaccine which confers protective immunity to both a Streptococcal infection and infection by another pathogenic agent is also within the scope of the invention.

20 As indicated above, the vaccines of the present invention are expected to elicit a protective immune response against infections caused by species and strains of *Streptococcus* other than strain of *S. pneumoniae* deposited with that ATCC.

25 Further within the scope of the invention are whole cell and whole viral vaccines. Such vaccines may be produced recombinantly and involve the expression of one or more of the *S. pneumoniae* polypeptides described in Table 1. For example, the *S. pneumoniae* polypeptides of the present invention may be either secreted or localized intracellular, on the cell surface, or in the periplasmic space. Further, when a recombinant virus is used, the *S.*



*pneumoniae* polypeptides of the present invention may, for example, be localized in the viral envelope, on the surface of the capsid, or internally within the capsid. Whole cells vaccines which employ cells expressing heterologous proteins are known in the art. See, e.g., Robinson, K. *et al.*, *Nature Biotech.* 15:653-657 (1997); Sirard, J. *et al.*, *Infect. Immun.* 65:2029-2033 (1997); Chabalgoity, J. *et al.*, *Infect. Immun.* 65:2402-2412 (1997). These cells may be administered live or may be killed prior to administration. Chabalgoity, J. *et al.*, *supra*, for example, report the successful use in mice of a live attenuated *Salmonella* vaccine strain which expresses a portion of a platyhelminth fatty acid-binding protein as a fusion protein on its cells surface.

A multi-component vaccine can also be prepared using techniques known in the art by combining one or more *S. pneumoniae* polypeptides of the present invention, or fragments thereof, with additional non-streptococcal components (e.g., diphtheria toxin or tetanus toxin, and/or other compounds known to elicit an immune response). Such vaccines are useful for eliciting protective immune responses to both members of the *Streptococcus* genus and non-streptococcal pathogenic agents.

The vaccines of the present invention also include DNA vaccines. DNA vaccines are currently being developed for a number of infectious diseases. Boyer, J *et al.*, *Nat. Med.* 3:526-532 (1997); reviewed in Spier, R., *Vaccine* 14:1285-1288 (1996). Such DNA vaccines contain a nucleotide sequence encoding one or more *S. pneumoniae* polypeptides of the present invention oriented in a manner that allows for expression of the subject polypeptide. The direct administration of plasmid DNA encoding *B. burgdorferi* OspA has been shown to elicit protective immunity in mice against borrelial challenge. Luke, C. *et al.*, *J. Infect. Dis.* 175:91-97 (1997).

The present invention also relates to the administration of a vaccine which is co-administered with a molecule capable of modulating immune responses. Kim, J. *et al.*, *Nature Biotech.* 15:641-646 (1997), for example, report the enhancement of immune responses produced by DNA immunizations when DNA sequences encoding molecules which stimulate the immune response are co-administered. In a similar fashion, the vaccines of the present invention may be co-administered with either nucleic acids encoding immune modulators or the immune modulators themselves. These immune modulators include granulocyte macrophage colony stimulating factor (GM-CSF) and CD86.

The vaccines of the present invention may be used to confer resistance to streptococcal infection by either passive or active immunization. When the

vaccines of the present invention are used to confer resistance to streptococcal infection through active immunization, a vaccine of the present invention is administered to an animal to elicit a protective immune response which either prevents or attenuates a streptococcal infection. When the vaccines of the present invention are used to confer resistance to streptococcal infection through passive immunization, the vaccine is provided to a host animal (*e.g.*, human, dog, or mouse), and the antisera elicited by this antisera is recovered and directly provided to a recipient suspected of having an infection caused by a member of the *Streptococcus* genus.

The ability to label antibodies, or fragments of antibodies, with toxin molecules provides an additional method for treating streptococcal infections when passive immunization is conducted. In this embodiment, antibodies, or fragments of antibodies, capable of recognizing the *S. pneumoniae* polypeptides disclosed herein, or fragments thereof, as well as other *Streptococcus* proteins, are labeled with toxin molecules prior to their administration to the patient. When such toxin derivatized antibodies bind to *Streptococcus* cells, toxin moieties will be localized to these cells and will cause their death.

The present invention thus concerns and provides a means for preventing or attenuating a streptococcal infection resulting from organisms which have antigens that are recognized and bound by antisera produced in response to the polypeptides of the present invention. As used herein, a vaccine is said to prevent or attenuate a disease if its administration to an animal results either in the total or partial attenuation (*i.e.*, suppression) of a symptom or condition of the disease, or in the total or partial immunity of the animal to the disease.

The administration of the vaccine (or the antisera which it elicits) may be for either a "prophylactic" or "therapeutic" purpose. When provided prophylactically, the compound(s) are provided in advance of any symptoms of streptococcal infection. The prophylactic administration of the compound(s) serves to prevent or attenuate any subsequent infection. When provided therapeutically, the compound(s) is provided upon or after the detection of symptoms which indicate that an animal may be infected with a member of the *Streptococcus* genus. The therapeutic administration of the compound(s) serves to attenuate any actual infection. Thus, the *S. pneumoniae* polypeptides, and fragments thereof, of the present invention may be provided either prior to the onset of infection (so as to prevent or attenuate an anticipated infection) or after the initiation of an actual infection.

5 The polypeptides of the invention, whether encoding a portion of a native protein or a functional derivative thereof, may be administered in pure form or may be coupled to a macromolecular carrier. Example of such carriers are proteins and carbohydrates. Suitable proteins which may act as macromolecular carrier for enhancing the immunogenicity of the polypeptides of the present invention include keyhole limpet hemacyanin (KLH) tetanus toxoid, pertussis toxin, bovine serum albumin, and ovalbumin. Methods for coupling the polypeptides of the present invention to such macromolecular carriers are disclosed in Harlow *et al.*, *Antibodies: A Laboratory Manual, 2nd Ed.*; Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York (1988), the entire disclosure of which is incorporated by reference herein.

10 A composition is said to be "pharmacologically acceptable" if its administration can be tolerated by a recipient animal and is otherwise suitable for administration to that animal. Such an agent is said to be administered in a "therapeutically effective amount" if the amount administered is physiologically significant. An agent is physiologically significant if its presence results in a detectable change in the physiology of a recipient patient.

15 While in all instances the vaccine of the present invention is administered as a pharmacologically acceptable compound, one skilled in the art would recognize that the composition of a pharmacologically acceptable compound varies with the animal to which it is administered. For example, a vaccine intended for human use will generally not be co-administered with Freund's adjuvant. Further, the level of purity of the *S. pneumoniae* polypeptides of the present invention will normally be higher when administered to a human than when administered to a non-human animal.

20 As would be understood by one of ordinary skill in the art, when the vaccine of the present invention is provided to an animal, it may be in a composition which may contain salts, buffers, adjuvants, or other substances which are desirable for improving the efficacy of the composition. Adjuvants are substances that can be used to specifically augment a specific immune response. These substances generally perform two functions: (1) they protect the antigen(s) from being rapidly catabolized after administration and (2) they nonspecifically stimulate immune responses.

25 Normally, the adjuvant and the composition are mixed prior to presentation to the immune system, or presented separately, but into the same site of the animal being immunized. Adjuvants can be loosely divided into several groups based upon their composition. These groups include oil adjuvants (for example, Freund's complete and incomplete), mineral salts (for

example,  $\text{AlK}(\text{SO}_4)_2$ ,  $\text{AlNa}(\text{SO}_4)_2$ ,  $\text{AlNH}_4(\text{SO}_4)$ , silica, kaolin, and carbon), polynucleotides (for example, poly IC and poly AU acids), and certain natural substances (for example, wax D from *Mycobacterium tuberculosis*, as well as substances found in *Corynebacterium parvum*, or *Bordetella pertussis*, and members of the genus *Brucella*. Other substances useful as adjuvants are the saponins such as, for example, Quil A. (Superfos A/S, Denmark). Preferred adjuvants for use in the present invention include aluminum salts, such as  $\text{AlK}(\text{SO}_4)_2$ ,  $\text{AlNa}(\text{SO}_4)_2$ , and  $\text{AlNH}_4(\text{SO}_4)$ . Examples of materials suitable for use in vaccine compositions are provided in *Remington's Pharmaceutical Sciences* (Osol, A, Ed, Mack Publishing Co, Easton, PA, pp. 1324-1341 (1980), which reference is incorporated herein by reference).

The therapeutic compositions of the present invention can be administered parenterally by injection, rapid infusion, nasopharyngeal absorption (intranasopharangeally), dermoabsorption, or orally. The compositions may alternatively be administered intramuscularly, or intravenously. Compositions for parenteral administration include sterile aqueous or non-aqueous solutions, suspensions, and emulsions. Examples of non-aqueous solvents are propylene glycol, polyethylene glycol, vegetable oils such as olive oil, and injectable organic esters such as ethyl oleate. Carriers or occlusive dressings can be used to increase skin permeability and enhance antigen absorption. Liquid dosage forms for oral administration may generally comprise a liposome solution containing the liquid dosage form. Suitable forms for suspending liposomes include emulsions, suspensions, solutions, syrups, and elixirs containing inert diluents commonly used in the art, such as purified water. Besides the inert diluents, such compositions can also include adjuvants, wetting agents, emulsifying and suspending agents, or sweetening, flavoring, or perfuming agents.

Therapeutic compositions of the present invention can also be administered in encapsulated form. For example, intranasal immunization of mice against *Bordetella pertussis* infection using vaccines encapsulated in biodegradable microsphere composed of poly(DL-lactide-co-glycolide) has been shown to stimulate protective immune responses. Shahin, R. *et al.*, *Infect. Immun.* 63:1195-1200 (1995). Similarly, orally administered encapsulated *Salmonella typhimurium* antigens have also been shown to elicit protective immunity in mice. Allaoui-Attarki, K. *et al.*, *Infect. Immun.* 65:853-857 (1997). Encapsulated vaccines of the present invention can be administered by

a variety of routes including those involving contacting the vaccine with mucous membranes (e.g., intranasally, intracolonicly, intraduodenally).

Many different techniques exist for the timing of the immunizations when a multiple administration regimen is utilized. It is possible to use the compositions of the invention more than once to increase the levels and diversities of expression of the immunoglobulin repertoire expressed by the immunized animal. Typically, if multiple immunizations are given, they will be given one to two months apart.

According to the present invention, an "effective amount" of a therapeutic composition is one which is sufficient to achieve a desired biological effect. Generally, the dosage needed to provide an effective amount of the composition will vary depending upon such factors as the animal's or human's age, condition, sex, and extent of disease, if any, and other variables which can be adjusted by one of ordinary skill in the art.

The antigenic preparations of the invention can be administered by either single or multiple dosages of an effective amount. Effective amounts of the compositions of the invention can vary from 0.01-1,000  $\mu\text{g/ml}$  per dose, more preferably 0.1-500  $\mu\text{g/ml}$  per dose, and most preferably 10-300  $\mu\text{g/ml}$  per dose.

Having now generally described the invention, the same will be more readily understood through reference to the following example which is provided by way of illustration, and is not intended to be limiting of the present invention, unless specified.

### **Examples**

#### ***Example 1: Expression and Purification of S. pneumoniae Polypeptides in E. coli***

The bacterial expression vector pQE10 (QIAGEN, Inc., 9259 Eton Avenue, Chatsworth, CA, 91311) is used in this example for cloning of the nucleotide sequences shown in Table 1 and for expressing the polypeptides identified in Table 1. The components of the pQE10 plasmid are arranged such that the inserted DNA sequence encoding a polypeptide of the present invention expresses the polypeptide with the six His residues (i.e., a "6 X His tag") covalently linked to the amino terminus.

The DNA sequences encoding the desired portions of the polypeptides of Table 1 are amplified using PCR oligonucleotide primers from either a DNA library constructed from *S. pneumoniae*, such as the one deposited by the inventors at the ATCC for convenience, ATCC Deposit No. 97755, or from

DNA isolated from the same organism such as the *S. pneumoniae* strain deposited with the ATCC as Deposit No. 55840. A list of PCR primers which can be used for this purpose is provided in Table 3, below. The PCR primers anneal to the nucleotide sequences encoding both the amino terminal and carboxy terminal amino acid sequences of the desired portion of the polypeptides of Table 1. Additional nucleotides containing restriction sites to facilitate cloning in the pQE10 vector were added to the 5' and 3' primer sequences, respectively. Such restriction sites are listed in Table 3 for each primer. In each case, the primer comprises, from the 5' end, 4 random nucleotides to prevent "breathing" during the annealing process, a restriction site (shown in Table 3), and approximately 15 nucleotides of *S. pneumoniae* ORF sequence (the complete sequence of each cloning primer is shown as SEQ ID NO:227 through SEQ ID NO:452).

For cloning the polypeptides of Table 1, the 5' and 3' primers were selected to amplify their respective nucleotide coding sequences. One of ordinary skill in the art would appreciate that the point in the protein coding sequence where the 5' primer begins may be varied to amplify a DNA segment encoding any desired portion of the complete amino acid sequences described in Table 1. Similarly, one of ordinary skill in the art would further appreciate that the point in the protein coding sequence where the 3' primer begins may also be varied to amplify a DNA segment encoding any desired portion of the complete amino acid sequences described in Table 1.

The amplified DNA fragment and the pQE10 vector are digested with the appropriate restriction enzyme(s) and the digested DNAs are then ligated together. The ligation mixture is transformed into competent *E. coli* cells using standard procedures such as those described in Sambrook *et al.*, *Molecular Cloning: a Laboratory Manual, 2nd Ed.*; Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y. (1989). Transformants are identified by their ability to grow under selective pressure on LB plates. Plasmid DNA is isolated from resistant colonies and the identity of the cloned DNA confirmed by restriction analysis, PCR and DNA sequencing.

Clones containing the desired constructs are grown overnight ("O/N") in liquid culture under selection. The O/N culture is used to inoculate a large culture, at a dilution of approximately 1:25 to 1:250. The cells are grown to an optical density at 600 nm ("OD600") of between 0.4 and 0.6. Isopropyl-b-D-thiogalactopyranoside ("IPTG") is then added to a final concentration of 1 mM to induce transcription from the *lac* repressor sensitive promoter, by inactivating

the *lacI* repressor. Cells subsequently are incubated further for 3 to 4 hours. Cells are then harvested by centrifugation.

The cells are stirred for 3-4 hours at 4 C in 6M guanidine-HCl, pH 8. The cell debris is removed by centrifugation, and the supernatant containing the protein of interest is loaded onto a nickel-nitrilo-tri-acetic acid ("NiNTA") affinity resin column (available from QIAGEN, Inc., *supra*). Proteins with a 6x His tag bind to the Ni-NTA resin with high affinity and can be purified in a simple one-step procedure (for details see: The QIAexpressionist, 1995, QIAGEN, Inc., *supra*). Briefly, the supernatant is loaded onto the column in 6 M guanidine-HCl, pH8, the column is first washed with 10 volumes of 6 M guanidine-HCl, pH8, then washed with 10 volumes of 6 M guanidine-HCl pH6, and finally the polypeptide is eluted with 6 M guanidine-HCl, pH 5.0.

The purified protein is then renatured by dialyzing it against phosphate-buffered saline (PBS) or 50 mM Na-acetate, pH 6 buffer plus 200 mM NaCl. Alternatively, the protein can be successfully refolded while immobilized on the Ni-NTA column. The recommended conditions are as follows: renature using a linear 6M-1M urea gradient in 500 mM NaCl, 20% glycerol, 20 mM Tris/HCl pH7.4, containing protease inhibitors. The renaturation should be performed over a period of 1.5 hours or more. After renaturation the proteins can be eluted by the addition of 250 mM imidazole. Imidazole is removed by a final dialyzing step against PBS or 50 mM sodium acetate pH6 buffer plus 200 mM NaCl. The purified protein is stored at 4°C or frozen at -80°C.

The DNA sequences encoding the amino acid sequences of Table 1 may also be cloned and expressed as fusion proteins by a protocol similar to that described directly above, wherein the pET-32b(+) vector (Novagen, 601 Science Drive, Madison, WI 53711) is preferentially used in place of pQE10.

Each of the polynucleotides shown in Table 1, was successfully amplified and subcloned into pQE10 as described above using the PCR primers shown in Table 3. These pQE10 plasmids containing the DNAs of Table 1, except SP023, SP042, SP054, SP063, SP081, SP092, SP114, SP122, SP123, SP126, and SP127, were deposited with the ATCC as a pooled deposit as a convenience to those of skill in the art. This pooled deposit was desposited on October 16, 1997 and given ATCC Deposit No. 209369. Those of ordinary skill in the art appreciate that isolating an individual plasmid from the pooled deposit is trivial provided the information and reagents described herein. Each of the deposited clones is capable of expressing its encoded *S. pneumoniae* polypeptide.

## Example 2: Immunization and Detection of Immune Responses

### Methods

#### Growth of bacterial inoculum, immunization of Mice and Challenge with *S pneumoniae*.

Propagation and storage of, and challenge by *S. pneumoniae* are performed essentially as described in Aaberge, I.S. et al., Virulence of *Streptococcus pneumoniae* in mice: a standardized method for preparation and frozen storage of the experimental bacterial inoculum, *Microbial Pathogenesis*, 18:141 (1995), incorporated herein by reference.

Briefly, Todd Hewitt (TH) broth (Difco laboratories, Detroit, MI) with 17% FCS, and horse blood agar plates are used for culturing the bacteria. Both broth and blood plates are incubated at 37°C in a 5% CO<sub>2</sub> atmosphere. Blood plates are incubated for 18 hr. The culture broth is regularly 10-fold serially diluted in TH broth kept at room temperature and bacterial suspensions are kept at room temperature until challenge of mice.

For active immunizations C3H/HeJ mice (The Jackson Laboratory, Bar Harbor, ME) are injected intraperitoneally (i.p.) at week 0 with 20 g of recombinant streptococcal protein, or phosphate-buffered saline (PBS), emulsified with complete Freund's adjuvant (CFA), given a similar booster immunization in incomplete Freund's adjuvant (IFA) at week 4, and challenged at week 6. For challenge *S. pneumoniae* are diluted in TH broth from exponentially-growing cultures and mice are injected subcutaneously (s.c.) at the base of the tail with 0.1 ml of these dilutions (serial dilutions are used to find medium infectious dose). Streptococci used for challenge are passaged fewer than six times *in vitro*. To assess infection, blood samples are obtained from the distal part of the lateral femoral vein into heparinized capillary tubes. A 25 ul blood sample is serially 10-fold diluted in TH broth, and 25 ul of diluted and undiluted blood is plated onto blood agar plates. The plates are incubated for 18 hr. and colonies are counted.

Other methods are known in the art, for example, see Langermann, S. et al., *J. Exp. Med.*, 180:2277 (1994), incorporated herein by reference.



### *Immunoassays*

Several immunoassay formats are used to quantify levels of streptococcal-specific antibodies (ELISA and immunoblot), and to evaluate the functional properties of these antibodies (growth inhibition assay). The ELISA and immunoblot assays are also used to detect and quantify antibodies elicited in response to streptococcal infection that react with specific streptococcal antigens. Where antibodies to certain streptococcal antigens are elicited by infection this is taken as evidence that the streptococcal proteins in question are expressed *in vivo*. Absence of infection-derived antibodies (seroconversion) following streptococcal challenge is evidence that infection is prevented or suppressed. The immunoblot assay is also used to ascertain whether antibodies raised against recombinant streptococcal antigens recognize a protein of similar size in extracts of whole streptococci. Where the natural protein is of similar, or identical, size in the immunoblot assay to the recombinant version of the same protein, this is taken as evidence that the recombinant protein is the product of a full-length clone of the respective gene.

### *Enzyme-Linked Immunosorbant Assay (ELISA).*

The ELISA is used to quantify levels of antibodies reactive with streptococcus antigens elicited in response to immunization with these streptococcal antigens. Wells of 96 well microtiter plates (Immunlon 4, Dynatech, Chantilly, Virginia, or equivalent) are coated with antigen by incubating 50  $\mu$ l of 1  $\mu$ g/ml protein antigen solution in a suitable buffer, typically 0.1 M sodium carbonate buffer at pH 9.6. After decanting unbound antigen, additional binding sites are blocked by incubating 100  $\mu$ l of 3% nonfat milk in wash buffer (PBS, 0.2% Tween 20, pH 7.4). After washing, duplicate serial two-fold dilutions of sera in PBS, Tween 20, 1% fetal bovine serum, are incubated for 1 hr, removed, wells are washed three times, and incubated with horseradish peroxidase-conjugated goat anti-mouse IgG. After three washes, bound antibodies are detected with H<sub>2</sub>O<sub>2</sub> and 2,2'-azino-di-(3-ethylbenzthiazoline sulfonate) (Schwan, T.G., *et al.*, *Proc. Natl. Acad. Sci. USA* 92:2909-2913 (1985)) (ABTS®, Kirkegaard & Perry Labs., Gaithersburg, MD) and A<sub>405</sub> is quantified with a Molecular Devices, Corp. (Menlo Park, California) Vmax™ plate reader. IgG levels twice the background level in serum from naive mice are assigned the minimum titer of 1:100.

***Sodiumdodecylsulfate-Polyacrylamide Gel Electrophoresis (SDS-PAGE) and Immunoblotting***

Using a single well format, total streptococcal protein extracts or recombinant streptococcal antigen are boiled in SDS/2-ME sample buffer before electrophoresis through 3% acrylamide stacking gels, and resolving gels of higher acrylamide concentration, typically 10-15% acrylamide monomer. Gels are electro-blotted to nitrocellulose membranes and lanes are probed with dilutions of antibody to be tested for reactivity with specific streptococcal antigens, followed by the appropriate secondary antibody-enzyme (horseradish peroxidase) conjugate. When it is desirable to confirm that the protein had transferred following electro-blotting, membranes are stained with Ponceau S. Immunoblot signals from bound antibodies are detected on x-ray film as chemiluminescence using ECL™ reagents (Amersham Corp., Arlington Heights, Illinois).

***Example 3: Detection of Streptococcus mRNA expression***

Northern blot analysis is carried out using methods described by, among others, Sambrook *et al.*, *supra*. to detect the expression of the *S. pneumoniae* nucleotide sequences of the present invention in animal tissues. A cDNA probe containing an entire nucleotide sequence shown in Table 1 is labeled with <sup>32</sup>P using the *rediprime*™ DNA labeling system (Amersham Life Science), according to manufacturer's instructions. After labeling, the probe is purified using a CHROMA SPIN-100™ column (Clontech Laboratories, Inc.), according to manufacturer's protocol number PT1200-1. The purified labeled probe is then used to detect the expression of *Streptococcus* mRNA in an animal tissue sample.

Animal tissues, such as blood or spinal fluid, are examined with the labeled probe using ExpressHyb™ hybridization solution (Clontech) according to manufacturer's protocol number PT1190-1. Following hybridization and washing, the blots are mounted and exposed to film at -70 C overnight, and films developed according to standard procedures.

It will be clear that the invention may be practiced otherwise than as particularly described in the foregoing description and examples.

Numerous modifications and variations of the present invention are possible in light of the above teachings and, therefore, are within the scope of the appended claims.

The entire disclosure of all publications (including patents, patent applications, journal articles, laboratory manuals, books, or other documents) cited herein are hereby incorporated by reference.

Table 1

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**SP001 nucleotide (SEQ ID NO:1)**

TAAAATCTACGACAATAAAAACTCAACTCATTGCTGACTTGGGTTCTGAACGCCGCGTCAATGCCCAAGC  
TAATGATATTCCACAGATTGTTAAGGCAATCGTTTCTATCGAAGACCATCGCTTCTTCGACCACAG  
GGGGATTGATACCATCCGTATCCTGGGAGCTTTCTTGCAGCAATCTGCAAAGCAATTCCCTCCAAGGTGG  
ATCAACTCTCACCCAACAGTTGATTAAGTTGACTTACTTTTCAACTTCGACTTCCGACCAGACTATTTTC  
TCGTAAGGCTCAGGAAGCTTGGTTAGCGATTTCAGTTAGAACAAAAAGCAACCAAGCAAGAAATCTTGAC  
CTACTATATAAATAAGGTCTACATGTCTAATGGGAACATGGAATGCAGACAGCAGCTCAAAACTACTA  
TGGTAAAGACCTCAATAATTTAAGTTTACCTCAGTTAGCCTTGCTGGCTGGAATGCCTCAGGCACCAAA  
CCAATATGACCCCTATTTCACATCCAGAAGCAGCCCAAGACCGCCGAACTTGGTCTTATCTGAAATGAA  
AAATCAAGGCTACATCTCTGCTGAACAGTATGAGAAAGCAGTCAATACACCAATTACTGATGGACTACA  
AAGTCTCAAATCAGCAAGTAATTACCCTGCTTACATGGATAATTACCTCAAGGAAGTCATCAATCAAGT  
TGAAGAAGAAACAGGCTATAACCTACTCACAACCTGGGATGGATGTCTACACAAATGTAGACCAAGAAGC  
TCAAAAACATCTGTGGGATATTTACAATACAGACGAATACGTTGCCATCCAGACGATGAATTGCAAGT  
CGCTTCTACCATTGTTGATGTTTCTAACGGTAAAGTCATTGCCAGCTAGGAGCACGCCATCAGTCAAG  
TAATGTTTCTCCTTGGAAATTAACCAAGCAGTAGAACAACACCGGACTGGGGATCAACTATGAAACCGAT  
CACAGACTATGCTCCTGCCTTGGAGTACGGTGTCTACGATTCAACTGCTACTATCGTTACAGATGAGCC  
CTATAACTACCCTGGGACAAATACTCCTGTTTATAACTGGGATAGGGGCTACTTTGGCAACATCACCTT  
GCAATACGCCCTGCAACAATCGCGAAACGTCCCAGCCGTGGAACTCTAAACAAGGTGGGACTCAACCG  
CGCCAAGACTTTCTTAAATGGTCTAGGAATCGACTACCCAAGTATTCACTACTCAAATGCCATTTCAAG  
TAACACAACCGAATCAGACAAAAAATATGGAGCAAGTAGTGAAGATGGCTGCTGCTTACGCTGCCCTT  
TGCAAATGGTGGAACTTACTATAAACCAATGTATATCCATAAAGTCGTCTTTAGTGATGGGAGTGAAAA  
AGAGTTCTCTAATGTGGAACTCGTGCCATGAAGGAAACGACAGCCTATATGATGACCGACATGATGAA  
AACAGTCTTGACTTATGGAACTGGACGAAATGCCTATCTTGCTTGGCTCCCTCAGGCTGGTAAAACAGG  
AACCTCTAACTATACAGACGAGGAAATTGAAAACACATCAAGACCTCTCAATTTGTAGCACCTGATGA  
ACTATTTGCTGGCTATACGCGTAAATATTCAATGGCTGTATGGACAGGCTATTCTAACCGTCTGACACC  
ACTTGTAGGCAATGGCCTTACGGTGCCTGCCAAAGTTTACCGCTCTATGATGACCTACCTGTCTGAAGG  
AAGCAATCCAGAAGATTGGAATATACCAGAGGGGCTCTACAGAAATGGAGAATTCGTATTTAAAAATGG  
TGCTCGTTCTACGTGGAACTCACCTGCTCCACAACAACCCCATCAACTGAAAGTTCAAGCTCATCATC  
AGATAGTTCAACTTCACAGTCTAGCTCAACCACTCCAAGCACAAATAATAGTACGACTACCAATCCTAA  
CAATAATACGCAACAATCAATACAACCCCTGATCAACAAAATCAGAATCCTCAACCAGCACAAACCA

**SP001 AMINO ACID (SEQ ID NO:2)**

KIYDNKNQLIADLGSERRVNAQANDIPTDLVKAIVSIEDHRFFDHRGIDTIRILGAFLRNLSNSLQGG  
STLTQQLIKLTYFSTSTSDQITSRKAQEAWLAIQLEQKATKQEILTYIYINKVYMSNGNYGMQTAAQNY  
GKDLNLSLPQLALLAGMPQAPNQYDPYSHPEAAQDRRLVLSEMNQGYISAEQYKAVNTPITDGLQ  
SLKSASNPAYMDNYLKEVINQVEETGYNLLTTGMDVYTNVDQEAQKHLWDIYNTDEYVAYPDDELQV  
ASTIVDVSNGKVIAQLGARHQSSNVSGINQAVETNRDWGSTMKPITDYAPALEYGVYDSTATIVHDEP  
YNYPGTNPVYNWDRGYFGNITLQYALQQSRNVPVAVETLNKVGLNRAKTFNLNGLIDYPSIHYSNAISS  
NTTESDKKYGASSEKMAAAYAAFANGGTYYKPMYIHKVVFSDGSEKEFSNVGTRAMKETTAAYMMTDMMK  
TVLTYGTGRNAYLAWLPQAGKTGTSNYTDEEIEENHIKTSQFVAPDELFAGYTRKYSMAVWVTGYSNRLTP  
LVGNGLTVAAKVYRSMMTYLSEGSNPEDWNIPEGLYRNGEFVKNGARSTWNSPAPQPPSTESSSSSS  
DSSTSQSSSTTPSTNNSTTNPNNNTQQSNTTPDQONQNPQPAQP

**SP004 nucleotide (SEQ ID NO:3)**

AAATTACAATACGGACTATGAATTGACCTCTGGAGAAAAATTACCTCTTCTTAAAGAGATTTAGGTTA  
CACTTATATTGGATATATCAAAGAGGGAAAAACGACTTCTGAGTCTGAAGTAAGTAATCAAAAGAGTTC  
AGTTGCCACTCCTACAAAACAACAAAAGGTGGATTATAATGTTACACCGAATTTTGTAGACCATCCATC  
AACAGTACAAGCTATTTCAGGAACAAACACCTGTTTCTTCAACTAAGCCGACAGAAGTTCAAGTAGTTGA  
AAAACCTTTCTCTACTGAATTAATCAATCCAAGAAAAGAAGAGAAACAATCTTCAGATTCTCAAGAACA  
ATTAGCCGAACATAAGAATCTAGAAACGAAGAAAGAGGAGAAGATTTCTCCAAAAGAAAAGACTGGGGT  
AAATACATTAAATCCACAGGATGAAGTTTATCAGGTCAATTGAACAAACCTGAACTCTTATATCGTGA  
GGAAACTATGGAGACAAAAATAGATTTTCAAGAAGAAATTCAAGAAAAATCCTGATTTAGCTGAAGGAAC  
TGTAAGAGTAAAACAAGAAGGTAAATTAGGTAAGAAAGTTGAAATCGTCAGAATATTCTCTGTAAACAA  
GGAAGAAGTTTCGCGAGAAATTGTTTCAACTTCAACGACTGCGCCTAGTCCAAGAATAGTCGAAAAAGG  
TACTAAAAAACTCAAGTTATAAAGGAACAACCTGAGACTGGTGTAGAACATAAGGACGTACAGTCTGG  
AGCTATTGTTGAACCCGAATTCAGCCTGAGTTGCCCCAAGCTGTAGTAAGTGACAAAGGCGAACCAGA  
AGTTCAACCTACATTACCCGAAGCAGTTGTGACCGACAAAGGTGAGACTGAGGTTCAACCAGAGCTCGCC  
AGATACTGTGGTAAGTGATAAAGGTGAACCAGAGCAGGTAGCACCGCTTCCAGAATATAAGGGTAATAT

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SP004 amino acid (SEQ ID NO:4)

SP006 nucleotide (SEQ ID NO:5)

TGAGAAATCAAGCTACACCCAAAGAGACTAGCGCTCAAAAGACAATCGTCCTTGCTACAGCTGGCGACGTT  
 GCCACCATTTGACTACGAAGACAAGGGCAATCTGACAGGCTTTGATATCGAAGTTTTAAAGGCAGTAGA  
 TGAAAAACTCAGCGACTACGAGATTCAATTCCAAAGAACCGCCTGGGAGAGCATCTTCCAGGACTTGA  
 TTCTGGTCACTATCAGGCTGCGGCCAATAACTTGAGTTACACAAAAGAGCGTGCTGAAAAATACCTTTA  
 CTCGCTTCCAATTTCCAACAATCCCCTCGTCCTTGTCAGCAACAAGAAAAATCCTTTGACTTCTCTTGA  
 CCAGATCGCTGGTAAAACAACACAAGAGGATACCGGAACTTCTAACGCTCAATTATCAATAAAGTGGAA  
 TCAGAAACACACTGATAATCCCGCTACAATTAATTTTCTGGTGAGGATATTGGTAAACGAATCCTAGA  
 CCTTGCTAACGGAGAGTTTGATTTCCTAGTTTTTGACAAGGTATCCGTTCAAAAGATTATCAAGGACCG  
 TGGTTTAGACCTCTCAGTCGTTGATTTACCTTTCTGCGAGTAGCCCCAGCAGTATATATCATCTTTCTCAAG  
 CGACCAAAAAGAGTTTAAAGAGCAATTTGATAAAGCGCTCAAAGAACTCTATCAAGACGGAACCCCTTGA  
 AAAAATCAGCAATACCTTACTTAGGTGGTTCTTACCTCCAGATCAATCTCAGTTACAA

SP006 amino acid (SEO ID NO:6)

ENQATPKETSQAQKTIVLATAGDVPPFDYEDKGNLTGFDIEVLKAVDEKLSDEYIQFRTAWESIFPGLD  
SGHYQAAANNLSYTKERAKEYLSLPISNNPLVLVSNKKNPLTSLDQIAGKTTQEDTGTNAQFINNWN  
QKHTDNPATINFSGEDIGKRILDLANGEFDLFLVDKVSQVKI IKDRGLDLSVVDLPSADSPSNYIIFSS  
DQKEFKEQFDKALKELYQDGTLEKLSNTYLGGSYLPDQSQLQ

SP007 nucleotide (SEQ ID NO:7)

TGTTAACCGCTCTTCTCGTAAACGACAGTTTCATCTTCTGATGTGAAGACAAAAGCAGCAATCGTCACTGA  
TACTGGTGGTGTTGATGACAAATCATTCAACCAATCAGCTTGGGAAGGTTTGCAGGCTTGGGGTAAAGA  
ACACAATCTTTCAAAGATAACGGTTTCACTTACTTCCAATCAACAAGTGAAGCTGACTACGCTAACAA  
CTTGCAACAAGCGGCTGGAAGTTACAACCTAATCTTCGGTGTTGGTTTTGCCCTTAATAATGCAGTTAA  
AGATGCAGCAAAAGAACACACTGACTTGAACATATGTCTTGATTGATGATGTGATTAAAGACCAAAAGAA  
TGTTGCGAGCGTAACCTTTCGCTGATAATGAGTCAGGTTACCTTGCAGGTGTGGCTGCAGCAAAAACAC  
TAAGACAAAACAAGTTGGTTTTGTAGGTGGTATCGAATCTGAAGTTATCTCTCGTTTTGAAGCAGGATT  
CAAGGCTGGTGTTGCGTCAGTAGACCCATCTATCAAAGTCCAAGTTGACTACGCTGGTTCATTGGTGA  
TGCGGCTAAAGGTAAAACAATTGCAGCCGCACAATACGCAGCCGGTGCAGATATTGTTTACCAAGTAGC  
TGGTGGTACAGGTGCAGGTGTCTTTGCAGAGGCAAAATCTCTCAACGAAAGCCGTCCTGAAAATGAAAA  
AGTTTGGGTATTCGGTGTTGATCGTGACCAAGAAGCAGAAGGTAAATACACTTCTAAAGATGGCAAAGA  
ATCAAACCTTTGTTCTTGATCTACTTTGAAACAAGTTGGTACAACCTGTAAAAGATATTTCTAACAAGGC  
AGAAAGAGGAGAATCCCTGGCGGCTCAAGTGATCGTTTACTCATTGAAGGATAAAGGGGTTGACTTGGC  
AGTACAAAACCTTTCAGAGGAAGGTAAAAAAGCTGTCGAAGATGCAAAAGCTAAAATCCTTGATGGAAG  
CGTAAAAAGTTCTGTAAAAA

Table 1

**SP007 amino acid (SEQ ID NO:8)**

GNRSSRNAASSSDVKTAAIVTDTGGVDDKSFNQSAWEGQLQAWGKEHNLSKDNGFTYFQSTSEADYANN  
LQQAAGSYNLI FGVGFALNNNAVKDAAKEHTDLNVLIDDVIKDQKNVASVTFADNESGYLAGVAAAKTT  
KTKQVGFVGGIESEVISRFEAGFKAGVASVDPSIKVQVDYAGSFGDAAKGKTIAAAQYAAGADIVYQVA  
GGTGAGVF AEAKSLNESRPENEKVWVIGVDRDQEAEGKYTSKDGKESNFVLVSTLKQVGTTVKDI SNKA  
ERGEFPFGQVIVYSLKDKGVDLAVTNLSEEGKKAVEDAKAKILDGSKVPEK

**SP008 nucleotide (SEQ ID NO:9)**

TGTGGAAATTTGACAGGTAACAGCAAAAAAGCTGCTGATTCAGGTGACAAACCTGTTATCAAAATGTAC  
CAAATCGGTGACAAACCAGACAACCTTGATGAATTGTTAGCAAATGCCAACAAATCATTGAAGAAAAA  
GTTGGTGCCAAATTTGGATATCCAATACCTTGGCTGGGGTGACTATGGTAAGAAAAATGTCAGTTATCACA  
TCATCTGGTGAAAACTATGATATTGCCCTTTCAGATAAATATATTGTAAATGCTCAAAAAGGTGCTTAC  
GCTGACTTGACAGAATTGTACAAAAAGGAAGGTAAAGACCTTTACAAAGCACTTGACCCAGCTTACATC  
AAGGGTAATACTGTAAATGGTAAGATTTACGCTGTTCCAGTTGCAGCCAACGTTGCATCATCTCAAAAC  
TTTGCCCTCAACGGAACCTCTCCTTGCTAAATATGGTATCGATATTTAGGTGTTACTTCTTACGAAACT  
CTTGAGCCAGTCTTGAAACAAATCAAAGAAAAAGCTCCAGACGTAGTACCATTGCTATTGGTAAAGTT  
TTCATCCCATCTGATAATTTTACTACCCAGTAGCAAACGGTCTTCCATTGCTTATCGACCTTGAAGGC  
GATACTACTAAAGTTGTAAACCGTTACGAAGTGCCTCGTTTCAAAGAACACTTGAAGACTCTTCACAAA  
TTCTATGAAGCTGGCTACATTCCAAAAGACGTGCAACAAGCGATACTTCCTTTGACCTTCAACAAGAT  
ACTTGGTTCTGTTCTGTAAGAAACAGTAGGACCAGCTGACTACGGTAACAGCTTGCTTTCACGTGTTGCC  
AACAAAGATATCCAAATCAAACCAATTACTAATTCATCAAGNAAAACCAAACACACAAGTTGCTAAC  
TTTGTCTATCTCAAAACACTCTAAGAACAAGAAAAATCAATGGAAATCTTGAACCTCTTGAATACGAAC  
CCAGAACTCTTGAACGGTCTTGTTTACGGTCCAGAAGGCAAGAACTGGGAAAAAATTGAAGGTAAAGAA  
AACCGTGTTCGCGTCTTGTATGGCTACAAAGGAAACACTCACATGGGTGGATGGAACACTGGTAACAAC  
TGGATCCTTTACATCAACGAAAACGTTACAGACCAACAAATCGAAAATTCTAAGAAAGAATTGGCAGAA  
GCTAAAGAATCTCCAGCGCTTGGATTATCTTCAATACTGACAATGTGAAATCTGAAATCTCAGCTATT  
GCTAACACAATGCAACAATTTGATTAACAGCTATCAACACTGGTACTGTAGACCCAGATAAAGCGATTCCA  
GAATTGATGAAAAAATTGAAATCTGAAGGTGCCTACGAAAAAGTATTGAACGAAATGCAAAAAACAATC  
GATGAATTCTTGA AAAACAAAAA

**SP008 amino acid (SEQ ID NO:10)**

CGNLTGNSKKAADSGDKPVIMYQIGDKPDNLDELLANANKIIIEKVGAKLDIQYLGWDYGGKMSVIT  
SSGENYDIAFADNYIVNAQKGAYADLTLYKKEGKDLYKALDPAYIKGNTVNGKIYAVPVAANVASSQN  
FAFNGTLLAKYGIDISGVTSYETLEPVLKQIKEKAPDVVPFAIGKVFIPSDNFDYPVANGLPFVIDLEG  
DTTKVVNRVEVPRFKEHLKTLHKFYEAGYIPKDVATSDTSFIDLQDFTWVRETVGPADYGNLSLLSRVA  
NKDIQIKPITNFIKXNQTTQVANFVISNNSKNKEKSMEILNLLNTNPELLNGLVYGPENKWEKIEGKE  
NRVRVLDGYKGNTHMGGWNTGNNWILYINENVTDQQIENSKKELAEAKESPALGFIFNTDNVSKSEISAI  
ANTMQQFDTAINTGTVPDKAIPELMEKLKSEGAYEKVLNEMQKQYDEFLKNKK

**SP009 nucleotide (SEQ ID NO:11)**

TGGTCAAGGAACTGCTTCTAAAGACAACAAAGAGGCAGAACTTAAGAAGGTTGACTTTATCCTAGACTG  
GACACCAAATACCAACCACACAGGGCTTTATGTTGCCAAGGAAAAAGGTTATTTCAAAGAAGCTGGAGT  
GGATGTTGATTTGAAATTGCCACCAGAAGAAAGTTCTTCTGACTTGTTATCAACGGAAGGCACCATT  
TGCAGTGTATTTCCAAGACTACATGGCTAAGAAATGGAAGAAAGGAGCAGGAATCACTGCCGTTGCAGC  
TATTGTTGAACACAATACATCAGGAATCATCTCTCGTAAATCTGATAATGTAAGCAGTCCAAAAGACTT  
GGTTGGTAAGAAATATGGGACATGGAATGACCAACTGAACTTGCTATGTTGAAAACCTTGGTAGAATC  
TCAAGGTGGAGACTTTGAGAAGGTTGAAAAAGTACCAAATAACGACTCAAACCTCAATCACACCGATTGC  
CAATGGCGCTCTTTGATACTGCTTGGATTTACTACGGTTGGGATGGTATCCTTGCTAAATCTCAAGGTGT  
AGATGCTAACTTCATGTACTTGAAAGACTATGTCAAGGAGTTTGACTACTATTACCAGTTATCATCGC  
AAACAACGACTATCTGAAAGATAACAAAGAAGAGCTCGCAAAGTCATCCAAGCCATCAAAAAGGCTA  
CCAATATGCCATGGAACATCCAGAAGAAGCTGCAGATATTCTCATCAAGAATGCACCTGAACTCAAGGA  
AAAACGTGACTTTGTCTCATCGAATCTCAAAAATACTTGTCAAAAAGAAATACGCAAGCGACAAGGAAAAATG  
GGGTCAATTTGACGCAGCTCGCTGGAATGCTTTCTACAAATGGGATAAAGAAAATGGTATCCTTAAAGA  
AGACTTGACAGACAAAGGCTTCACCAACGAATTTGTGAAA

**SP009 amino acid (SEQ ID NO:12)**

Table 1

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QGGTASKDNKEAELKKVDFILDWTPNTNHTGLYVAKEKGYFKEAGVDVDLKLPPPESSSDLVINGKAPF  
AVYFQDYMAKKLEKAGITAVAAIVEHNTSGIISRKSDNVSSPKDLVGKKYGTWNDPTELAMKTLIVES  
QGGDFEKVEKVPNNDSNSITPIANGVFDTAWIYYGWDGILAKSQGVDFANFMYLKDYVKEFDYYSPIVIA  
NNDYLDKNKEEARKVIQAIKKGYQYAMEHPPEEAADILIKNAPELKEKRDFVIESQKYLKEYASDKEKW  
QGFDAARWNAFYKWDKENGILKEDLTDKGFTNEFVK

**SP010 nucleotide (SEQ ID NO:13)**

TAGCTCAGGTGGAACGCTGGTTCATCCTCTGGAACAACTGCCAAAGCTCGCACTATCGATGAAAT  
CAAAAAAGCGGTGAAGTGCAGTTCGCCGTGTTTGGAGATAAAAAACCGTTTGGCTACGTTGACAATGA  
TGTTTCTACCAAGGTACGCTACGATATTGAAGTAGGGAACCACTAGCTCAAGACCTTGGTGTCAAGGT  
TAAATACATTTTACGTGATGCTGCCAACCGTGCGGAATACTTGATTTCAAACAAGGTAGATATTACTCT  
TGCTAACTTTACAGTAAGTACGACGAACGTAAAGAAACAAGTTGATTTTGCCTTCCATATATGAAAGTTTC  
TCTGGGTGTCGTATCACCTAAGACTGGTCTCATTACAGACGTCAAACAACCTGAAGGTAAACCTTAAT  
TGTCACAAAAGGAACGACTGCTGAGACTTATTTTGAAGAAGATCATCCAGAAATCAAACCTCAAAAAATA  
CGACCAATACAGTGACTCTTACCAAGCTCTTCTTGACGGACGTGGAGATGCCTTTTCAACTGACAATAC  
GGAAGTTCTAGCTTGGGCGCTTGAATAAAGGATTGAAAGTAGGAATTACTTCCCTCGGTGATCCCGA  
TACCATTGCGGCAGCAGTTCAAAAAGGCAACCAAGAATTGCTAGACTTCATCAATAAAGATATTGAAAA  
ATTAGGCAAGGAAAACCTTCTCCACAAGGCCTATGAAAAGACACTTCACCCAACCTACGGTGACGCTGC  
TAAAGCAGATGACCTGGTTGTTGAAGGTGGAAGTTGAT

**SP010 amino acid (SEQ ID NO:14)**

SSGGNAGSSSGKTTAKARTIDEIKKSGELRIAVFGDKKPGYVDNDGSTKVRDYDIELGNQLAQDLGVKV  
KYISVDAANRAEYLISNKVDITLANFTVTDERKKQVDFALPYMKVSLGVVSPKTGLITDVKQLEGKTLI  
VTKGTTAETYFEKNHPEIKLKQYDQYSDSYQALLDGRGDAFSTDNTEVLAWALENKGFEVGITSLGDPD  
TIAAAVQKGNQELDLFINKDIEKLGKENFFHKAYEKTLLHPTYGDAAKADDLVVEGGKVD

**SP011 nucleotide (SEQ ID NO:15)**

CTCCAACATATGGTAAATCTGCGGATGGCACAGTGACCATCGAGTATTTCAACCAGAAAAAAGAAATGAC  
CAAAACCTTGGAAAGAAATCACTCGTGATTTTGAAGAAGGAAAACCTAAGATCAAGGTCAAAGTCGTCAA  
TGTACCAAATGCTGGTGAAGTATTGAAGACACGCGTTCTCGCAGGAGATGTGCCTGATGTGGTCAATAT  
TTACCCACAGTCCATCGAACTGCAAGAATGGGCAAAAGCAGGTGTTTTTGAAGATTGAGCAACAAAGA  
CTACCTGAAACGCGTGAAAAATGGCTACGCTGAAAAATATGCTGTAAACGAAAAAGTTTACAACGTTCC  
TTTTACAGCTAATGCTTATGGAATTTACTACAACAAAGATAAATTCGAAGAAGTGGGCTTGAAGGTCC  
TGAAACCTGGGATGAATTTGAACAGTTAGTCAAAGATATCGTTGCTAAAGGACAAACACCATTTGGAAT  
TGCAGGTGCAGATGCTTGGACACTCAATGGTTACAATCAATTAGCCTTTGCGACAGCAACAGGTGGAGG  
AAAAGAAGCAAATCAATACCTTCGTTATTCTCAACCAATGCCATTAAATTGTCCGATCCGATTATGAA  
AGATGATATCAAGGTCATGGACATCCTTCGCATCAATGGATCTAAGCAAAAGAACTGGGAAGGTGCTGG  
CTATACCGATGTTATCGGAGCCTTCGCACGTGGGGATGTCTCATGACACCAATGGGTCTTGGGCGAT  
CACAGCGATTAATGAACAAAAACCGAACTTAAAGATTGGGACCTTCATGATTCCAGGAAAAAGAAAAAGG  
ACAAAGCTTAACCGTTGGTGGCGGAGACTTGGCATGGTCTATCTCAGCCACCACCAACATCCAAAAGA  
AGCCAATGCCTTTGTGGAATATATGACCCGTCCAGAAGTCATGCAAAAATACTACGATGTGGACGGATC  
TCCAACAGCGATCGAAGGGGTCAAACAAGCAGGAGAAGATTACCGCTTGCTGGTATGACCGAATATGC  
CTTTACGGATCGTCACTTGGTCTGGTTGCAACAATACTGGACCAAGTGAAGCAGACTTCCATACCTTGAC  
CATGAACATATGCTTGGACCGGTGATAAACAAGGCATGGTCAATGATTTGAATGCCTTCTTTAACCCGAT  
GAAAGCGGATGTGGAT

**SP011 amino acid (SEQ ID NO:16)**

SNYKGSADGVTITIEYFNQKKEMTKLEEITRDFEKENPKIKVKVNVNPNAGEVLKTRVLGADVPDVVNI  
YPQSIELQEWAKAGVFEDLSNKDYLRVKNGYAEKYAVNEKVYNVPFTANAYGIYYNKDKFEELGLKVP  
ETWDEFEQLVKDIVAKGQTPFGIAGADAWTLNGYNQLAFATATGGGKEANQYLYRSQPNAILKLSDPIMK  
DDIKVMDILRINGSKQKNWEGAGYTDVIGAFARGDVLMTPNGSWAITAINEQKPNFKIGTFMIPGKEKG  
QSLTVGAGDLAWSISATTKHPKEANAFVEYMTREVMQKYYDVGDSPTAIEGVKQAGEDSPLAGMTEYA  
FTDRHLVWLQQYWTSEADFHTLTMNYVLTGDKQGMVNDLNAFFNPMKADVD

**SP012 nucleotide (SEQ ID NO:17)**

TGGGAAAAATTCTAGCGAAACTAGTGGAGATAATTGGTCAAAGTACCAGTCTAACAAGTCTATTACTAT  
TGGATTTGATAGTACTTTTGTTCGAATGGGATTTGCTCAGAAAGATGGTTCTTATGCAGGATTTGATAT  
TGATTTAGCTACAGCTGTTTTTGAAAAATACGGAATCACGGTAAATTGGCAACCGATTGATTGGGATTT

Table 1

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GAAAGAAGCTGAATTGACAAAAGGAACGATTGATCTGATTGGAATGGCTATTCCGCTACAGACGAACG  
CCGTGAAAAGTGCGTTTTCAGTAACATCATATATGAAGAATGAGCAGGTATTGGTTACGAAGAAATCATC  
TGGTATCACGACTGCAAAGGATATGACTGGAAAGACATTAGGAGCTCAAGCTGGTTCATCTGGTTATGC  
GGACTTTGAAGCAAATCCAGAAATTTTGAAGAATATTGTCGCTAATAAGGAAGCGAATCAATACCAAAC  
CTTTAATGAAGCCTTGATTGATTTGAAAAACGATCGAATTGATGGTCTATTGATTGACCGTGTCTATGC  
AACTATTATTATTAGAAGCAGAAGGTGTTTTAAACGATTATAATGTCTTTACAGTTGGACTAGAAACAGA  
AGCTTTTGCGGTTGGAGCCCGTAAGGAAGATACAACTTGGTTAAGAAGATAAATGAAGCTTTTTCTAG  
TCTTTACAAGGACGGCAAGTTCGAAGAAATCAGCCAAAAATGGTTTGGAGAAGATGTAGCAACCAAAGA  
AGTAAAAGAAGGACAG

**SP012 nucleotide (SEQ ID NO:18)**

GKNSSETSGDNWSKYQSNKSITIGFDSTFVPMGFAQKDGSYAGFDIDLATAVFEKYGITVNWQPIDWDL  
KEAELTKGTIDLIWNGYSATDERREKVAFSNSYMKNEQVLVTKKSSGITTAKDMTGKTLGAQAGSSGYA  
DFEANPEILKNIVANKEANQYQTFNEALIDLKNDRIDGLLIDRVYANYYLEAEGVLNDYNVFTVGLETE  
AFAVGARKEDTNLVKKINEAFSSLYKDGKFQEISQKWFGEDEVATKEVKEGQ

**SP013 nucleotide (SEQ ID NO:19)**

TGCTAGCGGAAAAAAGATACAACTTCTGGTCAAAAACTAAAAGTTGTTGCTACAACTCAATCATCGC  
TGATATTACTAAAAATATTGCTGGTGACAAAATTGACCTTCATAGTATCGTTCCGATTGGGCAAGACCC  
ACACGAATACGAACCACTTCTGAAGACGTTAAGAAAACCTCTGAGGCTAATTTGATTTTCTATAACGG  
TATCAACCTTGAAACAGGTGGCAATGCTTGGTTTACAAAATTGGTAGAAAATGCCAAGAAAACCTGAAAA  
CAAAGACTACTTCCGAGTCAGCGACGGCGTTGATGTTATCTACCTTGAAGGTCAAAATGAAAAAGGAAA  
AGAAGACCCACACGCTTGGCTTAACCTTGAAAACGGTATTATTTTTGCTAAAAATATCGCCAAACAATT  
GAGCGCCAAAGACCTTAACAATAAAGAATTCTATGAAAAAATCTCAAAGAATATACTGATAAGTTAGA  
CAAACCTTGATAAAGAAAGTAAGGATAAATTTAATAAGATCCCTGCTGAAAAGAACTCATTGTAACCAG  
CGAAGGAGCATTCAAATACTTCTCTAAAGCCTATGGTGTCCCAAGTGCTTACATCTGGGAAATCAATAC  
TGAAGAAGAAGGAACCTCTGAACAAATCAAGACCTTGGTTGAAAAACTTCGCCAAACAAAAGTTCATC  
ACTCTTTGTAGAATCAAGTGTGGATGACCGTCCAATGAAAACTGTTTCTCAAGACACAAACATCCCAAT  
CTACGCTCAAATCTTTACTGACTCTATCGCAGAACCAAGGTAAAGAAGGCGACAGCTACTACAGCATGAT  
GAAATACAACCTTGACAAGATTGCTGAAGGATTGGCAAAA

**SP013 amino acid (SEQ ID NO:20)**

ASGKKDTSQQLKVVATNSIIADITKNIAGDKIDLHSIVPIGQDPHEYEPLPEDVKKTSEANLIFYNG  
INLETGGNAWFTKLVENAKKTENKDYFAVSDGVDVIYLEGQNEKGKEDPHAWLNLENGIIFAKNIAKQL  
SAKDPNNKEFYENLKEYTDKLDKLDKESKDKFNKIPAEKKLIVTSEGAFKYFSKAYGVPSAYIWEINT  
EEEGTPEQIKTLVEKLRQTKVPSLFEVSSVDDRPMTVSQDTNPIPIYAQIFTDSIAEQKEGDSYYSMM  
KYNLDKIAEGLAK

**SP014 nucleotide (SEQ ID NO:21)**

TGGCTCAAAAAATACAGCTTCAAGTCCAGATTATAAGTTGGAAGGTGTAACATTCCCGCTTCAAGAAAA  
GAAAACATTGAAGTTTATGACAGCCAGTTCACCGTTATCTCCTAAAGACCCAAATGAAAAGTTAATTTT  
GCAACGTTTGGAGAAGGAACTGGCGTTTATATTGACTGGACCACTACCAATCCGACTTTGCAGAAAA  
ACGTAACCTGGATATTTCTAGTGGTGATTTACCAGATGCTATCCACAACGACGGAGCTTCAGATGTGGA  
CTTGATGAACCTGGGCTAAAAAAGGTGTTATTATTCCAGTTGAAGATTTGATTGATAAATACATGCCAAA  
TCTTAAGAAAAATTTGGATGAGAAACCAGAGTACAAGGCCCTTGATGACAGCACCTGATGGGCACATTTA  
CTCATTTCCATGGATTGAAGAGCTTGGAGATGGTAAAGAGTCTATTACAGTGTCAACGATATGGCTTG  
GATTAACAAAGATTGGCTTAAGAACTTGGTCTTGAAATGCCAAAACTACTGATGATTTGATTAAAGT  
CCTAGAAGCTTTCAAAAACGGGGATCCAAATGGAAATGGAGAGGCTGATGAAATTCCATTTTCATTTAT  
TAGTGGTAACGGAACGAAGATTTTAAATTCTTATTGCTGCATTTGGTATAGGGGATAACGATGATCA  
TTTAGTAGTAGGAAATGATGGCAAAGTTGACTTCACAGCAGATAACGATAACTATAAAGAAGGTGTCAA  
ATTTATCCGCTCAATTGCAAGAAAAAGGCCTGATTGATAAAGAAGCTTTCGAACATGATTGGAATAGTTA  
CATTGCTAAAGGTCATGATCAGAAATTTGGTGTCTTACTTTACATGGGATAAGAATAATGTTACTGGAAG  
TAACGAAAGTTATGATGTTTTACCAGTACTTGTGCGACCAAGTGGTCAAAAACACGTAGCTCGTACAAA  
CGGTATGGGATTTGCACGTGACAAGATGGTTATTACCAGTGTAACAAAAACCTAGAATTGACAGCTAA  
ATGGATTGATGCACAATACGCTCCACTCCAATCTGTGCAAAATAACTGGGGAACCTACGGAGATGACAA  
ACAACAAAACATCTTTGAATTGGATCAAGCGTCAAAATAGTCTAAAACACTTACCCTAAACGGAACCTGC  
ACCAGCAGAACTTCGTCAAAAGACTGAAGTAGGAGGACCACTAGCTATCCTAGATTACATCTATGGTAA  
AGTAACAACCATGCTGATGATGCCAAATGGCGTTTGGATCTTATCAAAGAATATTATGTTCTTACAT



Table 1

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GAGCAATGTCAATAACTATCCAAGAGTCTTTATGACACAGGAAGATTTGGACAAGATTGCCCATATCGA  
AGCAGATATGAATGACTATATCTACCGTAAACGTGCTGAATGGATTGTAAATGGCAATATTGATACTGA  
GTGGGATGATTACAAGAAAGAACTTGAAAAATACGGACTTTCTGATTACCTCGCTATTAAACAAAAATA  
CTACGACCAATACCAAGCAAACAAAAAC

**SP014 amino acid (SEQ ID NO:22)**

GSKNTASSPDYKLEGVTFPLQEKKTLKFM TASSPLSPKDPNEKLILQRLEKETGVHIDWTNYQSDFAEK  
RNLDISSGDLPD A IHNDGASDV DLMNWAKKGVII PVEDLIDKYPNLKKILDEKPEYKALMTAPDGHYI  
SFPWIEELGDGKESIHSVNDMAWINKDWLKKLGLEMPKTTDDL IKVLEAFKNGDPNGNGEAD EIPFSFI  
SGNGNEDFKFLFAAFGIGDNDHLLVVGNDGKVDFTADNDNYKEGVKFI RQLQEKGLIDKEAF EHDWNSY  
IAKGHDQKFGVYFTW DKNVNTGSNESYDVL PVLAGPSGQKHVARTNGMGFARDKMVITSVNKNLELTAK  
WIDAQYAPLQSVQNNWGTYGDDKQONIFELDQASNSLKHPLPLNGTAPAE LRQKTEVGGPLAILDSYYGK  
VTTMPDDAKWRLLDIKEYYVPYMSNVNNYPRVFM TQEDLDKIAHIEADMNDYIYRKRAEWIVNGNIDTE  
WDDYKKELEKYGLSDYLAIKQKYDQYQANKN

**SP015 nucleotide (SEQ ID NO:23)**

TAGTACAAACTCAAGCACTAGTCAGACAGAGACCAGTAGCTCTGCTCCAACAGAGGTAACCATTA AAAAG  
TTC ACTGGACGAGGTCAAAC TTTCCAAAGTTCCTGAAAAGATTGTGACCTTTGACCTCGGCGCTGCGGA  
TACTATTCGCGCTTTAGGATTTGAAAAAATATCGTCGGAATGCCTACAAAAACTGTTCCGACTTATCT  
AAAAGACCTAGTGGGAAGTGTCAAAAATGTTGGTCTCTATGAAAGAACCTGATTTAGAAGCTATCGCCGC  
CCTTGAGCCTGATTTGATTATCGCTTCGCCACGTACACAAAAATTCGTAGACAAATTCAAAGAAATCGC  
CCCAACCGTTCTCTTCCAAGCAAGCAAGGACGACTACTGGACTTCTACCAAGGCTAATATCGAATCCTT  
AGCAAGTGCCCTTCGGCGAAAAGTGGTACACAGAAAGCCAAGGAAGAATTGACCAAGCTAGACAAGAGCAT  
CCAAGAAGTCGCTACTAAAAATGAAAGCTCTGACAAAAAGCCCTTGCGATCCTCTTAATGAAGGAAA  
AATGGCAGCCTTTGGTGCCAAATCTCGTTTCTCTTTCTGTACCAAACCTTGAAATTCAAACCAACTGA  
TACAAAATTTGAAGACTCACGCCACGGACAAGAAGTCAGCTTTGAAAGTGTCAAAGAAATCAACCCTGA  
CATCCTCTTTGTCAACCGTACCTTGCCATCGGTGGGACAACCTCTAGCAACGACGGTGTCTCTAGA  
AAATGCCCTTATCGCTGAAACACCTGCTGCTAAAAATGGTAAGATTATCCAAC TAACACCGACCTCTG  
GTATCTAAGCGGAGCGCGGACTTGAATCAACAAAACCTCATGATTGAAGACATACAAAAAGCTTTGAAA

**SP015 amino acid (SEQ ID NO:24)**

STNSSTSQTETSSSAPTEVTIKSSLDEVKLSKVPEKIVTFDLGAADTIRALGF EKNIVGMPTKTVP TYL  
KDLVGTVKNVSGMKEPDLEAIAALEPDLIIASPR TQKFVDKFK EIAPTVLFQASKDDYWTSTKANIESL  
ASAFGETGTQAKEELTKLDKSIQEVATKNES S DKKALAILLNEGKMAAFGA KS RFSFLYQTLKFKPTD  
TKFEDSRHQEVSFESVKEINPDILFVINRTLAIGD N SSNDGVLENALIAETPAAKNGKIIQLTPDLW  
YLSGGGLESTKLMIEDIQKALK

**SP016 nucleotide (SEQ ID NO:25)**

TGGCAATTC TGGCGGAAGTAAAGATGCTGCCAAATCAGGTGGTGACGGTGCCAAAACAGAAATCACTTG  
GTGGGCATTCCAGTATTTACCCAAGAAAAAAGTGGTGACGGTGTTGGAACCTTATGAAAAATCAATCAT  
CGAAGCGTTTGAAAAAGCAAACCCAGATATAAAAGTGAAATTGGAACCATCGACTTCAAGTCAGGTCC  
TGAAAAAATCACAAACAGCCATCGAAGCAGGAACAGCTCCAGACGTACTCTTTGATGCACCAGGACGTAT  
CATCCAATACGGTAAAAACGGTAAATTGGCTGAGTTGAATGACCTCTTCACAGATGAATTTGTTAAAGA  
TGTCACAATGAAAACATCGTACAAGCAAGTAAAGCTGGAGACAAGGCTTATATGTATCCGATTAGTTC  
TGCCCCATTCTACATGGCAATGAACAAGAAAATGTTAGAAGATGCTGGAGTAGCAAACCTTGTA AAAAGA  
AGGTTGGACAAGTATGATTTTGAAAAAGTATTGAAAGCACTTAAAGACAAGGGTTACACACCAGGTTC  
ATTGTTTCAGTTCTGGTCAAGGGGGAGACCAAGGAACACGTGCCTTTATCTCTAACCTTTATAGCGGTTT  
TGTAACAGATGAAAAAGTTAGCAAATATACAAGTATGATGATCTAAATTCGTCAAAGGTCTTGAAAAAGC  
AACTAGCTGGATTAAAGACAATTTGATCAATAATGGTTTCAATTTGACGGTGGGGCAGATATCCAAAA  
CTTTGCCAACGGTCAAACATCTTACACAATCCTTTGGGCACCAGCTCAAAATGGTATCCAAGCTAAACT  
TTTAGAAGCAAGTAAGGTAGAAGTGGTAGAAGTACCATTCCCATCAGACGAAGGTAAGCCAGCTCTTGA  
GTACCTTGTAACCGGTTTGCAGTATTCAACAATAAAGACGACAAGAAAGTCGCTGCATCTAAGAAAT  
CATCCAGTTTATCGCAGATGACAAGGAGTGGGGACCTAAAGACGTAGTTTCGTACAGGTGCTTTCCAGT  
CCGTACTTCATTTGGAAGAACTTTATGAAGACAAACGCATGGAACAATCAGCGGCTGGACTCAATACTA  
CTCACCATACTACAACACTATTGATGGATTTGCTGAAATGAGAACACTTTGGTTCCCAATGTTGCAATC  
TGTATCAAATGGTGACGAAAAACAGCAGATGCTTTGAAAGCCTTCACTGAAAAAGCGAACGAAACAAT  
CAAAAAAGCTATGAAACAA

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**SP016 amino acid (SEQ ID NO:26)**

GNSGGSKDAAKSGGDGAKTEITWWAFPVFTQEKTDGVDGVTYEKSIIEAFEKANPDIKVKLETIDFKSGP  
EKITTAIEAGTAPDVLFDAPGRIIQYGKNGKLAELNDLFTDEFVKDVNNENIVQASKAGDKAYMPISS  
APFYMANMKMLEADAGVANLVKEGWTTDDFEKVLKALKDKGYTPGSLFSSGQGGDQGTRAFISNLYSGS  
VTDEKVSQYTTDDPKFVKGLEKATSWIKDNLINNGSQFDGGADIQNFANGQTSYITLWAPAQNGIQAKL  
LEASKVEVVEVPFSPDEGKPALEYLVNGFAVFNKDDKVAASKKFIQFIADDDKEWGPDKDVVRTGAFPV  
RTSFGKLYEDKRMETISGWTQYYSPYNTIDGFAEMRTLWFPMLQSVSNGDEKPADALKAFTEKANETI  
KKAMKQ

**SP017 nucleotide (SEQ ID NO:27)**

TTCACAAGAAAAACAAAAATGAAGATGGAGAACTAAGACAGAACAGACAGCCAAAGCTGATGGAAC  
AGTCGGTAGTAAGTCTCAAGGAGCTGCCAGAGAAAGCAGAAGTGGTCAATAAAGGTGATTACTACAG  
CATTCAAGGGAAATACGATGAAATCATCGTAGCCAAACAAACACTATCCATTGTCTAAAGACTATAATCC  
AGGGGAAAAATCCAACAGCCAAGGCAGAGTTGGTCAAACCTCATCAAAGCGATGCAAGAGGCAGGTTTCCC  
TATTAGTGATCATTACAGTGGTTTTAGAACTTATGAAACTCAGACCAAGCTCTATCAAGATTATGTCAA  
CCAAGATGGAAGGCAGCAGCTGACCGTTACTCTGCCCCGTCTGGCTATAGCGAACACAGACAGGCTT  
GGCCTTTGATGTGATTGGGACTGATGGTGATTGGTGACAGAAGAAAAAGCAGCCCAATGGCTCTTGGA  
TCATGCAGCTGATTATGGCTTTGTTGTCCGTTATCTCAAAGGCAAGGAAAAGGAAACAGGCTATATGGC  
TGAAGAAATGGCACCTGCGTTATGTAGGAAAAGAAGCTAAAGAAATTGCTGCAAGTGGTCTCAGTTTGA  
AGAATACTATGGCTTTGAAGGCGGAGACTACGTCGAT

**SP017 amino acid (SEQ ID NO:28)**

SQEKTKNEDGETKTEQTAKADGTVGSKSQGAAQKKAEEVVKNGDYYSIQGKYDEIIVANKHYPLSKDYNP  
GENPTAKAELVKLIKAMQEAGFPISDHYSGFRSYETQTKLYQDYVNQDGKAAADRY SARPGYSEHQTL  
AFDVIQDGLVTEEKAAQWLLDHAADYGFVVRYLKGKEKETGYMAEEWHLRYVGKEAKEIAASGLSLE  
EYYGFEGGDYVD

**SP019 nucleotide (SEQ ID NO:29)**

GAAAGGTCTGTGGTCAAATAATCTTACCTGCGGTTATGATGAAAAATAATCTTGGAAAAATATAAATAT  
AAAAATACCTGAAGAAAAAATATCAGTTATTATTGGGTCAAATGGTTGTGGGAAATCAACACTCATTA  
AACCTTGCTCTGACTTATAAAGCCATTAGAGGGAGAAGTATTGCTTGATAATAAATCAATTAATTCTTA  
TAAAGAAAAAGATTTAGCAAAACACATAGCTATATTACCTCAATCTCCAATAATCCCTGAATCAATAAC  
AGTAGCTGATCTTGTAAGCCGTGGTCGTTTCCCTACAGAAAGCCTTTTAAGAGTCTTGAAAAGATGA  
CCTTGAAATAATAACAGATCAATGGTTAAGGCCAATGTTGAAGATCTAGCAAATAACCTAGTTGAAGA  
ACTTTCTGGGGTCAAAGGCAAAGAGTATGGATAGCTCTAGCCCTAGCCCAAGATACAAGTATCCTACT  
TTTAGATGAGCCAACTACTTACTTGGATATCTCATATCAAATAGAACTATTAGACCTCTTGACTGATCT  
AAACCAAAAATATAAGACAACCATTTGCATGATTTTGCACGATATAAATCTAACAGCAAGATACGCTGA  
TTACCTATTTGCAATTAAAGAAGGTAACTTGTTCAGAGGGAAAGCCTGAAGATATACTAAATGATAA  
ACTAGTTAAAGATATCTTTAATCTTGAAGCAAAAATTATACGTGACCCTATTTCCAATTCGCCTCTAAT  
GATTCCTATTGGCAAGCACCATGTTAACCT

**SP019 amino acid (SEQ ID NO:30)**

KGLWSNNLTCGYDEKIILENINIKIPEEKISVIIGSNGCGKSTLIKTLRLIKPLEGEVLLDNKSINSY  
KEKDLAKHIAILPQSPPIIPESITVADLVSRGRFPYRKPFKSLGKDDLEIINRSMVKANVEDLANNLVEE  
LSGGQRQVRWIALALAQDTSILLLEPTTYLDISYQIELLDLLTDLNQKYKTTICMILHDINLTARYAD  
YLFAIKEGKLVAEGKPEDILNDKLVKIDIFNLEAKIIRDPISNSPLMIPIGKHVS

**SP020 nucleotide (SEQ ID NO:31)**

AAACTCAGAAAAGAAAGCAGACAATGCAACAATATCAAAATCGCAACTGTAAACCGTAGCGGTTCTGA  
AGAAAAACGTTGGGACAAAATCCAAGAATTGGTTAAAAAAGACGGAATTACCTTGGAATTTACAGAGTT  
CACAGACTACTCACAACCAAAACAAAGCAACTGCTGATGGCGAAGTAGATTTGAACGCTTTCCAACACTA  
TAACTTCTTGAACAACTGGAACAAAGAAAACGGAAAAGACCTTGTAGCGATTGCAGATACTTACATCTC  
TCCAATCCGCCCTTACTCAGGTTTGAATGGAAGTGCCAACAAGTACACTAAAGTAGAAGACATCCCAGC  
AAACGGAGAAATCGCTGTACCGAATGACGCTACAAACGAAAGCCGTGCGCTTTATTTGCTTCAATCAGC  
TGGCTTGATTAAATTTGATGTTTCTGGAAGTCTTGTCAACAGTTGCCAACATCAAAGAAAATCCAAA  
GAAGTTGAAAATCACTGAATTTGACGCTAGCCAAACAGCTCGTTTATTGTCATCAGTTGACGCTGCCGT  
TGTAACAATAACCTTTCGTTACAGAAGCAAAATTGGACTACAAGAAATCACTTTTCAAAGAACAAGCTGA  
TGAAACTCAAACAATGGTACAACATCATTTGTGCAAAAAAAGATTGGGAAACATCACCTAAGGCTGA

Table 1

TGCTATCAAGAAAGTAATCGCAGCTTACCACACAGATGACGTGAAAAAAGTTATCGAAGAATCATCAGA  
TGGTTTGGATCAACCAGTTTGG

**SP020 amino acid (SEQ ID NO:32)**

NSEKKADNATTIKIATVNRSGSEEKRWDKIQELVKKDGITLEFTFTDYSQPNKATADGEVDLNAFQHY  
NFLNNWNKENGKDLVAIADTYISPIRLYSGLNGSANKYTKVEDIPANGEIAPNDATNESRALYLLQSA  
GLIKLDVSGTALATVANIKENPKNLKITELDASQTARSLSSVDAAVVNNTFVTEAKLDYKKSFLKEQAD  
ENSKQWYNIIVAKKDWETSPKADAIAAYHTDDVKVIEESSDGLDQPVW

**SP021 nucleotide (SEQ ID NO:33)**

TTCGAAAGGGTCAGAAGGTGCAGACCTTATCAGCATGAAAGGGGATGTCATTACAGAACATCAATTTTA  
TGAGCAAGTGAAAAGCAACCTTCAGCCCAACAAGTCTTGTTAAATATGACCATCCAAAAAGTTTGTGA  
AAAACAATATGGCTCAGAGCTTGATGATAAAGAGGTTGATGATACTATTGCCGAAGAAAAAACAATA  
TGGCGAAAACCTACCAACGTGTCTTGTCAACAAGCAGGTATGACTCTTGAAACACGTAAAGCTCAAATTCG  
TACAAGTAAATTAGTTGAGTTGGCAGTTAAGAAGGTAGCAGAAGCTGAATTGACAGATGAAGCCTATAA  
GAAAGCCTTTGATGAGTACACTCCAGATGTAACGGCTCAAATCATCCGTCTTAATAATGAAGATAAGGC  
CAAAGAAGTTCTCGAAAAAGCCAAGGCAGAAGGTGCTGATTTTGCTCAATTAGCCAAAGATAATTCAAC  
TGATGAAAAAACAAGAAAAATGGTGGAGAAATTACCTTTGATTCTGCTTCAACAGAAGTACCTGGAGC  
AAGTCCAAAAAAGCCGCTTTTCGCTTTTAGATGTGGGATGGTGTCTTGGATGTGGATTACAGCAACTG  
GGGCACACCAAGCCTACAG

**SP021 amino acid (SEQ ID NO:34)**

SKGSEGADLISMKGDVITEHQFYEQVKSNPQAQVLLNMTIQKVFEEKQYGESELDDEKVVDDTIAEEKQY  
GENYQRVLSQAGMTLETRKAQIRTSKLVELAVKKVAEAEELTDEAYKKAFDEYTPDVTAQIIRLNEDKA  
KEVLEKAKAEGADFAQLAKDNSTDEKTKENGGEITFDSASTEVPGASPKKPLFAFRCGMVFLDVDYSNW  
GTPSLQ

**SP022 nucleotide (SEQ ID NO:35)**

GGGGATGGCAGCTTTTAAAAATCCTAACAATCAATACAAAGCTATTACAATTGCTCAAACCTCTAGGTGA  
TGATGCTTCTTCAGAGGAATTGGCTGGTAGATATGGTTCTGCTGTTCAGTGTACAGAAGTGACTGCCTC  
AAACCTTTCAACAGTTAAACTAAAGCTACGGTTGTAGAAAAACCACTGAAAGATTTTAGAGCGTCTAC  
GTCTGATCAGTCTGGTTGGGTGGAATCTAATGGTAAATGGTATTTCTATGAGTCTGGTGATGTGAAGAC  
AGGTTGGGTGAAAACAGATGGTAAATGGTACTATTTGAATGACTTAGGTGTCATGCAGACTGGATTTGT  
AAAATTTTCTGGTACCTGGTATTACTTGAGCAATTCAGGTGCTATGTTTACAGGCTGGGGAACAGATGG  
TAGCAGATGGTTCTACTTTGACGGCTCAGGAGCTATGAAGACAGGCTGGTACAAGGAAAATGGCACTTG  
GTATTACCTTGACGAAGCAGGTATCATGAAGACAGGTTGGTTTAAAGTCGGACCACACTGGTACTATGC  
CTACGGTTCAGGAGCTTTGGCTGTGAGCACAACAACACCAGATGGTTACCGTGTAAATGGTAATGGTGA  
ATGGGTAAAC

**SP022 amino acid (SEQ ID NO:36)**

GMAAFKNPNQYKAITIAQTLGDDASSEELAGRYGSAVQCTEVTASNLSVTKTKATVVEKPLKDFRST  
SDQSGWVESNGKWYFYESGDVKTGWVKTDGKWYYLNDLGVMTGFVKFSGSWYYLSNSGAMFTGWGTDG  
SRWYFDGSGAMKTGWYKENGWYLLDEAGIMKTGWFKVGPWHYYAYGSGALAVSTTTPDGYRVNGNE  
WVN

**SP023 nucleotide (SEQ ID NO:37)**

AGACGAGCAAAAAATTAAGCAAGCAGAAGCGGAAGTTGAGAGTAAACAAGCTGAGGCTACAAGGTAAAA  
AAAAATCAAGACAGATCGTGAAGAAGCAGAAGAAGCTAAACGAAGAGCAGATGCTAAAGAGCAAGG  
TAAACCAAGGGGCGGGCAAAACGAGGAGTTCTTGGAGAGCTAGCAACACCTGATAAAAAAGAAAATGA  
TGCGAAGTCTTCAGATTCTAGCGTAGGTGAAGAACTCTTCCAAGCCCATCCCTGAAACCAGAAAAAAA  
GGTAGCAGAAGCTGAGAAGAAGGTTGAAGAAGCTAAGAAAAAAGCCGAGGATCAAAAAGAAGAAGATCG  
CCGTAACCTACCCAACCAATACTTACAAAACGCTTGAACCTTGAAATTGCTGAGTCCGATGTGGAAGTTAA  
AAAAGCGGAGCTTGAACCTAGTAAAAGAGGAAGCTAAGGAACCTCGAAACGAGGAAAAAGTTAAGCAAGC  
AAAAGCGGAAGTTGAGAGTAAAAAGCTGAGGCTACAAGGTTAGAAAAATCAAGACAGATCGTAAAAA  
AGCAGAAGAAGAAGCTAAACGAAAAGCAGCAGAAGAAGATAAAGTTAAAGAAAAACCAGCTGAACAACC  
ACAACCAGCGCCGGCTCCAAAAGCAGAAAAACCAGCTCCAGCTCCAAAACCAGAGAATCCAGCTGAACA  
ACCAAAAGCAGAAAAACCAGCTGATCAACAAGCTGAAGAAGACTATGCTCGTAGATCAGAAGAAGAATA  
TAATCGCTTGACTCAACAGCAACCGCCAAAAACTGAAAAACCAGCACACCATCTACTCCAAAAACAGG

Table 1

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CTGGAACAAGAAAACGGTATGTGGTACTTCTACAATACTGATGGTTCATGGCGACAGGATGGCTCCA  
AAACAATGGCTCATGGTACTACCTCAACAGCAATGGCGCTATGGCGACAGGATGGCTCCAAAACAATGG  
TTCATGGTACTATCTAAACGCTAATGGTTCATGGCAACAGGATGGCTCCAAAACAATGGTTCATGGTA  
CTACCTAAACGCTAATGGTTCATGGCGACAGGATGGCTCCAATACAATGGCTCATGGTACTACCTAAA  
CGCTAATGGTTCATGGCGACAGGATGGCTCCAATACAATGGCTCATGGTACTACCTAAACGCTAATGG  
TGATATGGCGACAGGTTGGGTGAAAGATGGAGATACCTGGTACTATCTTGAAGCATCAGGTGCTATGAA  
AGCAAGCCAATGGTTCAAAGTATCAGATAAATGGTACTATGTCAATGGCTCAGGTGCCCTTGAGTCAA  
CACAACCTGTAGATGGCTATGGAGTCAATGCCAATGGTGAATGGGTAAAC

**SP023 amino acid (SEQ ID NO:38)**

DEQKIKQAEAEVESKQAEATRLKKIKTDREEAEKEAKRRADAKEQKPKGRAKRGVPGELATPDKKEND  
AKSSDSSVGEETLPSPLKPEKKVAEAEKKVEEAKKAEDQKEEDRRNYPTNTYKTLELEIAESDVEVK  
KALELVKEEAKEPRNEEKVKQAKAEVESKKAETRLKIKTDKKAEEEEAKRKAEEEDKVKEKPAEQP  
QPAPAPKAEPAPAPKPNPAEQPKAEKPADQQAEDYARRSEEEYNRLTQQQPPKTEKPAQPSTPKTG  
WKQENGMWYFYNTDGSMATGWLQNNGSWYYLNSNGAMATGWLQNNGSWYYLNNANGSMATGWLQNNGSWY  
YLNANGSMATGWLQYNGSWYYLNNANGSMATGWLQYNGSWYYLNNANGDMATGWVKDGDWYYLEASGAMK  
ASQWFKVSDKWYYVNGSGALAVNTTVDGYGVNANGEWVN

**SP025 nucleotide (SEQ ID NO:39)**

CTGTGGTGAGGAAGAACTAAAAAGACTCAAGCAGCACAACAGCCAAAACAACAAACGACTGTACAACA  
AATTGCTGTTGGAAAAGATGCTCCAGACTTCACATTGCAATCCATGGATGGCAAAGAAGTTAAGTTATC  
TGATTTTAAGGGTAAAAAGGTTTACTTGAAGTTTTGGGCTTCATGGTGTGGTCCATGCAAGAAAAGTAT  
GCCAGAGTTGATGGAAGTAGCGGCGAAACCAGATCGTGATTTGCAAATTCCTACTGTCAATGCACCAGG  
AATTCAAGGTGAAAAAAGTGTGAGCAATTCACCAATGGTTCAGGAACAAGGATATAAGGATATCCC  
AGTTCTTTATGATACCAAAGCAACCACTTCCAAGCTTATCAAATTCGAAGCATTCCTACAGAATATT

**SP025 amino acid (SEQ ID NO:40)**

CGBEETKKTQAAQPKQQTTVQQIAVGKDAPDFTLQSMGKEVKLSDFKGGKVVYLFKFWASWCGPCKKSM  
PELMELAAKPDRDFEILTVIAPGIQGEKTVEQFPQWFQEQGYKDI PVLYDTKATTSKLIKFEAFQLNI

**SP028 nucleotide (SEQ ID NO:41)**

GACTTTTAAACAATAAACTATTGAAGAGTTGCACAATCTCCTTGTCTCTAAGGAAATTTCTGCAACAGA  
ATTGACCCAAGCAACACTTGAAAATATCAAGTCTCGTGAGGAAGCCCTCAATTCATTTGTACCATCGC  
TGAGGAGCAAGCTCTTGTCAAGCTAAAGCCATTGATGAAGCTGGAATTGATGCTGACAATGTCTTTTC  
AGGAATTCACCTTGCTGTAAAGGATAACATCTCTACAGACGGTATTCTCACAACCTGCTGCCTCAAAAAT  
GCTCTACAACCTATGAGCCAATCTTTGATGCGACagCTgTTGCCAATGCAAAAACCAAGGGCATGATTGT  
CGTTGGAAAGACCAACATGACGAATTTGCTATGGGTGGTTCAGGtGAAACTTCACACTACGGAGCAAC  
TAAAAACGCTTGGAACCACAGCAAGGTTCTGTGGTGGGTTCATCAAGTGGTTCGCGCAGCTGTAGCCTC  
AGGACAAGTTCGCTTGTCACCTTGGTTCGTGATACTGGTGGTTCATCCGCCAACCTGCTGCCTTCAACGG  
AATCGTTGGTCTCAAAACCAACCTACGGAACAGTTTCACGTTTCGGTCTCATTGCCTTTGGTAGCTCATT  
AGACCAGATTGGACCTTTTGCTCCTACTGTTAAGGAAAATGCCCTCTTGCTCAACGCTATTGCCAGCGA  
AGATGCTAAAGACTCTACTTCTGCTCCTGTCCGCATCGCCGACTTTACTTCAAAAATCGGCCAAGACAT  
CAAGGGTATGAAAATCGCTTTGCCTAAGGAATACCTAGGCGAAGGAATTGATCCAGAGGTAAAGGAAAC  
AATCTTAAACGCGGCCAAACACTTTGAAAAATTTGGGTGCTATCGTCGAAGAAGTCAGCCTTCTCACTC  
TAAATACGGTGTGGCGTTTATTACATCATCGCTTCATCAGAAGCTTCATCAAACCTGCAACGCTTCGA  
CGGTATCCGTTACGGCTATCGCGCAGAAGATGCAACCAACCTTGATGAAATCTATGTAAACAGCCGAAG  
CCAAGGTTTTGGTGAAGAGGTAAAACGTCGTATCATGCTGGGTACTTTCAAGTCTTTCAATCAGGTTACTA  
TGATGCCCTACTACAAAAAGGCTGGTCAAGTCCGTACCCTCATCATTCAAGATTTGAAAAAGTCTTCGC  
GGATTACGATTTGATTTTGGGTCCAACCTGCTCCAAGTGTGCTTATGACTTGGATTCTCTCAACCATGA  
CCCAGTTGCCATGTACTTAGCCGACCTATTGACCATACCTGTAAACTTGGCAGGACTGCCTGGAATTTTC  
GATTCCTGCTGGATTCTCTCAAGGTCTACCTGTGCGACTCCAATTGATTGGTCCCAAGTACTCTGAGGA  
AACCATTTACCAAGCTGCTGCTGCTTTTGAAGCAACAACAGACTACCACAAACAACAACCCGTGATTTT  
TGGAGGTGACAAC

**SP028 amino acid (SEQ ID NO:42)**

TFNNKTIEELHNLVLSKEISATELTQATLENIKSREEALNSFVTIAEEQALVQAKAIDEAGIDADNVLS  
GIPLAVKDNISTDGILTTAASKMLYNYEPIFDATAVANAKTKGMI VVGKTNMDEFAMGSGGETSHYGAT  
KNAWNHSHKVPGGSSSGSAAAVASGQVRLSLGSDTGGSI RQPAFNGIVGLKPTYGTVSRFGLIAFGSSL

Table 1

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DQIGPFAPTVKENALLLNIAISEDADKSTAPVRIADFTSKIGQDIKGMKIALPKEYLLEGIDPEVKET  
ILNAAKHFEKLGAIVEEVSLPHSKYGVAVYIIASSEASSNLQRFDGIRYGYRAEDATNLDEIYVNSRS  
QGFGEVKKRIMLGTFSLSSGYDAYKKAGQVRTLIQDFEKVFADYDLILGPTAPSVAYDLDLSLNHD  
PVAMYLADLLTIPVNLAGLPGISIPAGFSQGLPVGLQLIGPKYSEETIYQAAAAFEATTDYHKQQPVI  
GGDN

**SP030 nucleotide (SEQ ID NO:43)**

CTTTACAGGTAAACAACACTACAAGTCGGCGACAAGGCGCTTGATTTTCTCTTACTACAACAGATCTTTC  
TAAAAAATCTCTGGCTGATTTTGGATGGCAAGAAAAAGTCTTGAGTGTCGTTCTTCTATCGATACAGG  
CATCTGCTCAACTCAAACACGTCGTTTTAATGAAGAAITGGCTGGACTGGACAACACGGTCGTATTGAC  
TGTTTCAATGGACCTACCTTTTGCTCAAAAACGTTGGTGCGGTGCTGAAGGCCTTGACAATGCCATTAT  
GCTTTAGACTACTTTGACCATTCTTTGCGGCGCGATTATGCCCTCTTGATCAACGAATGGCACCTATT  
AGCACGCGCAGTCTTTGTCCTCGATACTGACAATACGATTGCTACGTTGAATACGTGGATAATATCAA  
TTCTGAGCCAAACTTCGAA

**SP030 amino acid (SEQ ID NO:44)**

FTGKQLQVGDKALDFSLTTTDLSSKSLADFDGKKVLSVVPSIDTGICSTQTRRFNEELAGLDNTVVL  
VSMDLPLFAQKRWCGAEGLDNAIMLSDFDHSFGRDYALLINEWHLLARAVFVLDTDNTIRYVEYVDNIN  
SEPNE

**SP031 nucleotide (SEQ ID NO:45)**

CCAGGCTGATACAAGTATCGCAGACATTCAAAAAAGAGGCGAACTGGTTGTGCGGTGTCAAACAAGACGT  
TCCCAATTTTGGTTACAAGATCCCAAGACCGGTACTTATTCTGGTATCGAAACCGACTTGGCCAAGAT  
GGTAGCTGATGAACCAAGGTCAAGATTCGCTATGTGCCGGTTACAGCACAAACCGCGGCCCCCTTCT  
AGACAATGAACAGGTGATATGGATATCGCGACCTTTACCATCACGGACGAACGCAAAAACTCTACAA  
CTTTACCAGTCCCTACTACACAGACGCTTCTGGATTTTGGTCAATAAATCTGCCAAATCAAAAAGAT  
TGAGGACCTAAACGGCAAAACCATCGGAGTCGCCCAAGGTTCTATACCCAACGCCTGATTACTGAACT  
GGGTAAAAAGAAAGGTCTGAAGTTTAAATTCGTCGAATCTGGTTCTTACCCAGAATTGATTACTTCCCT  
GCACGCTCATCGTATCGATACCTTTTCCGTTGACGCTCTATTCTATCTGGCTACACTAGTAAACGGAC  
AGCACTACTAGATGATAGTTTCAAGCCATCTGACTACGGTATTGTTACCAAGAAATCAAATACAGAGCT  
CAACGACTATCTTGATAACTTGGTTACTAAATGGAGCAAGGATGGTAGTTTGAGAACTTTATGACCG  
TTACAAGCTCAAACCATCTAGCCATACTGCAGAT

**SP031 amino acid (SEQ ID NO:46)**

QADTSIADIQKRGELVVGKQDVPNFGYXDPKGTGYSIETDLAKMVADELKVKIRYVPVTAQTRGPLL  
DNEQVDMDIATFTITDERKKLYNFTSPYYTDASGFLVNKSAKIKKIEDLNGKTIGVAQGSITQRLITEL  
GKKKGLKFKFVELGSYPELITSLHAHRIDTFVSDRSILSGYTSKRTALLDDSFKPSDYGIVTKKSNTL  
NDYLDNLVTKWSKDGSLQKLYDRYKLPSSHTAD

**SP032 nucleotide (SEQ ID NO:47)**

GTCTGTATCATTTGAAAAACAAAGAAACAAACCGTGGTGTCTTgACTTTCACTATCTCTCAAGACCAAAT  
CAAACCAGAATTGGACCGTGTCTTCAAGtCAGTGAAGAAATCTCTTAATGTTCCAGGTTTCCGTAAAGG  
TCACCTTCCACGCCCTATCTTCGACCAAAAATTTGGTGAAGAAGCTCTTTATCAAGATGCAATGAACGC  
ACTTTTGCCAAAACGCTTATGAAGCAGCTGTAAAAGAAGCTGGTCTTGAAGTGGTTGCCCAACCAAAAAT  
TGACGTAACCTCAATGGAAAAAGGTCAAGACTGGGTTATCACTGCTGAAGTCGTTACAAAACCTGAAGT  
AAAATTGGGTGACTACAAAACCTTGAAGTATCAGTTGATGTAGAAAAAGAAGTAAGTACGCTGATGT  
CGAAGAGCGTATCGAACGCGAACGCAACACCTGGCTGAATTGGTTATCAAGGAAGCTGCTGCTGAAAA  
CGGCGACACTGTTGTGATCGACTTCGTTGGTTCTATCGACGGTGTGAAATTTGACGGTGGAAGGTTGA  
AACTTCTCACTTGGACTTGGTTCAAGTCAATTCATCCCTGGTTTCGAAGACCAATTGGTAGGTCACTC  
AGCTGGCGAAACCGTTGATGTTATCGTAACATTCCCAGAAGACTACCAAGCAGAAGACCTTGCAGGTAA  
AGAAGCTAAATTCGTGACAACCTATCCACGAAGTAAAGCTAAAGAAGTCCGGCTCTTGACGATGAACT  
TGCAAAAGACATTGATGAAGAAGTTGAAACACTTGCTGACTTGAAAGAAAAATACAGCAAAGAATTGGC  
TGCTGCTAAAGAAGAAGCTTACAAAGATGCAGTTGAAGGTGCAGCAATTGATACAGCTGTAGAAAATGC  
TGAAATCGTAGAACTTCCAGAAGAAATGATCCATGAAGAAGTTCACCGTTCAGTAAATGAATTCCTTGG  
GAATTTGCAACGTCAAGGGATCAACCCTGACATGTACTTCCAAATCACTGGAAGTACTCAAGAAGACCT  
TCACAACCAATACCAAGCAGAAGCTGAGTCACGTACTAAGACTAACCTTGTATCGAAGCAGTTGCCAA  
AGCTGAAGGATTGATGCTTCAGAAGAAGAAATCCAAAAAGAAGTTGAGCAATTGGCAGCAGACTACAA

Table 1

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CATGGAAGTTGCACAAGTTCAAACTTGCTTTCAGCTGACATGTTGAAACATGATATCACTATCAAAAA  
AGCTGTTGAATTGATCACAAGCACAGCAACAGTAAAA

**SP032 amino acid (SEQ ID NO:48)**

SVSFENKETNRGVLTFTISQDIKPELDRVFKSVKSLNVPGFGRKGHLPRPIFDQKFGEEALYQDAMNA  
LLPNAYEAAVKEAGLEVVAQPKIDVTSMEKGQDWVITAENVTKPEVKLGDKNLEVSVDVEKEVTDADV  
EERIERERNLAEELVIKEAAAENGDTVVIDFVGSIDGVEFDGGKGENFSLGLGSGQFIPGFEDQLVGHS  
AGETVDVIVTTFPEDYQAEDLAGKEAKFVTTIHEVKAKEVPALDDELAKDIDEEVETLADLKEKYSKELA  
AAKEEAYKDAVEGAAIDTAVENAEIVELPEEMIHEEVHRSVNEFLGNLQRQGINPDMYFQITGTTQEDL  
HNQYQAEAESRTKTNLVIEAVAKAEGFDASEEEIQKEVEQLAADYNMEVAQVQNLLSADMLKHDITIKK  
AVELITSTATVK

**SP033 nucleotide (SEQ ID NO:49)**

TGGTCAAAAGGAAAGTCAGACAGGAAAGGGGATGAAAATTGTGACCAGTTTTTATCCTATCTACGCTAT  
GGTTAAGGAAGTATCTGGTGACTTGAATGATGTTCCGGATGATTCAGTCAAGTAGTGGTATTCACCTCTT  
TGAACCTTCGGCAAATGATATCGCAGCCATCTATGATGCAGATGTCTTTGTTTACCATTCTCATACT  
CGAATCTTGGGCAGGAAGTCTGGATCCAAATCTAAAAAATCCAAAGTGAAGGTCTTAGAGGCTTCTGA  
GGGAATGACCTTGAACGTGTCCCTGGACTAGAGGATGTGGAAGCAGGGGATGGAGTTGATGAAAAAAC  
GCTCTATGACCCTCACACATGGCTAGATCTGAAAAAGCTGGAGAAGAAGCCCAAATTATCGCTGATAA  
ACTTTTCAGAGGTGGATAGTGAGCATAAAGAGACTTATCAAAAAAATGCGCAACCTTTATCAAAAAAGCT  
CAGGAAT

**SP033 amino acid (SEQ ID NO:50)**

GQKESQTKGGMKIVTSFYPIYAMVKEVSGDLNDVRMIQSSSGIHSFEPSANDIAAIYDADVFVYHSHTL  
ESWAGSLDPNLKSKVKVLEASEGMTLERVPGLEDVEAGDGVDEKTLYPDHTWLDPEKAGEEAQIIADK  
LSEVDSEHKETYQKNAQPLSKLRLN

**SP034 nucleotide (SEQ ID NO:51)**

GAAGGATAGATATATTTTAGCATTGAGACATCCTGTGATGAGACCAGTGTGCGCGTCTTGAAAAACGA  
CGATGAGCTCTTGTCCAATGTCATTGCTAGTCAAATTGAGAGTCACAAACGTTTTGGTGGCGTAGTGCC  
CGAAGTAGCCAGTCGTCACCATGTCGAGGTCATTACAGCCTGTATCGAGGAGGCATTGGCAGAACGAGG  
GATTACCGAAGAGGACGTGACAGCTGTTGCGGTTACCTACGGACCAGGCTTGGTTCGGAGCCTTGCTAGT  
TGGTTTGTGAGCTGCCAAGGCCTTTGCTTGGGCTCACGGACTTCCACTGATTCTGTTAATCACATGGC  
TGGGCACCTCATGGCAGCTCAGAGTGTGGAGCCTTTGGAGTTTCCCTTGCTAGCCCTCTTGGTCAGCGG  
CGGACACACAGAGTTGGTTTATGTTTCGGAGGCAGGAGATTATAAGATTGTTGGGGAAACCCGTGATGA  
TGCGGTTGGTGAGGCTTATGATAAGGTCGCGCCGTGTCATGGGCTTGACCTATCCTGCAGGTCGTGAGAT  
TGACGAGCTGGCTCATCAGGGGCAGGATATTTATGATTTCCCCCGTGCCATGATTAAGGAAGATAATCT  
GGAGTTCTCCTTCTCAGGTTTGAAATCTGCCTTTATCAATCTTCATCACAATGCCGAGCAAAAGGGAGA  
AAGCCTGTCTACAGAAGATTGTGTGCTTCTTCCAAGCAGCAGTTATGGACATTCTCATGGCAAAAAC  
CAAGAAGGCTTTGGAGAAATATCCTGTTAAATCCTAGTTGTGGCAGGTGGTGTGGCAGCCAATAAAGG  
TCTCAGAGAACGCCTAGCAGCCGAAATCACAGATGTCAAGGTTATCATCCCCCTCTGCGACTCTGCGG  
AGACAATGCAGGTATGATTGCCTATGCCAGCGTCAGCNAGTGAACAAAGAAAACCTTCGAGGCTGGGA  
CCTCAATGCCAAACCAAGTCTTGCCCTTGATACCATGGAA

**SP034 amino acid (SEQ ID NO:52)**

KDRYILAFETSCDETSVAVLKNDDELLSNVIASQIESHKRFGGVVPEVASRHHVEVITACIEEALAEAG  
ITEEDVTAVAVTYGPGLVGALLVGLSAAKAFAWAHLPLIPVNHMAGHLMAAQSVLEPFLALLLVSG  
GHTELVYVSEAGDYKIVGETRDDAVGEAYDKVGRVMGLTYPAGREIDELAHQGDYDFPRAMIKEDNL  
EFSFSGLKSAFINLHHNAEQKGESLSTEDLCASFQAAVMDILMAKTKKALEKYPVKILVVAGGVAANKG  
LRERLAAEITDVKVIIPPLRLCGDNAGMIAYASVSXWNKENFAGWDLNAKPSLAFDTME

**SP035 nucleotide (SEQ ID NO:53)**

GGTAGTTAAAGTTGGTATTAACGGTTTCGGACGTATCGGTCGTCTTGCTTTCCGTCGTATCCAAAACGT  
AGAAGGTGTTGAAGTTACACGCATCAACGACCTTACAGATCCAGTTATGCTTGACACTTGTGAAATA  
CGACACAACCTCAAGGTCGTTTCGACGGTACTGTTGAAGTTAAAGAAGGTGGATTTGAAGTTAACGGTAA  
ATTCATCAAAGTTTCTGCTGAACGTGATCCAGAACAAATCGACTGGGCTACTGACGGTGTAGAAATCGT  
TCTTGAAGCTACTGGTTTCTTTGCTAAGAAAGAAGCAGCTGAAAAACACCTTAAAGGTGGAGCTAAAAA

Table 1

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AGTTGTTATCACTGCTCCTGGTGGAAACGACGTTAAAAACAGTTGTATTCAAACTAACCACGACGTTCT  
TGACGGTACTGAAACAGTTATCTCAGGTGCTTCATGTACTACAACTGCTTGGCTCCAATGGCTAAAGC  
TCTTCAAGACAACCTTTGGTGTGTGAAGGATTGATGACTACTATCCACGCTTACACTGGTGACCAAAT  
GATCCTTGACGGACCACACCGTGGTGGTGACCTTCGCCGCTGCTCGCGCTGGTGTGCAAACATCGTTCC  
TAACCTCAACTGGTGTGCAAAAGCTATCGGTCTTGTAATCCCAGAATTGAATGGTAAACTTGACGGATC  
TGCACAACGCGTTCCAACCTCAACTGGATCAGTTACTGAATTGGTAGCAGTTCTTGAAAAGAAGCTTAC  
TGTTGATGAAGTGAACGCAGCTATGAAAGCAGCTTCAAACGAATCATACGGTTACACAGAAGATCCAAT  
CGTATCTTCAGATATCGTAGGTATGTCTTACGGTTCATTGTTTGACGCAACTCAAACATAAGTTCTTGA  
CGTTGACGGTAAACAATTGGTTAAAGTTGTATCATGGTACGACAACGAAATGTCATACACTGCACAACCT  
TGTTTCGTACTCTTGAATACTTCGCAAAAATTGC

**SP035 amino acid (SEQ ID NO:54)**

VVKVINGFGRIGRLAFRIQNVGEVETRINDLTPVMLAHLKDYDTTQGRFDGTVEVKEGGFEVNGK  
FIKVSERDPEQIDWATDGVIEIVLEATGFFAKKEAAEKHLKGGAKKVITAPGGNDVKTVVFNTNHDVL  
DGTETVISGASCTTNCLAPMAKALQDNFVVEGLMTTIHAYTGDQMILDGPHRGDLRRARAGAANIVP  
NSTGAAKAIGLVIPELNGKLDGSAQRVPTPTGSVTELVAVLEKNVTVDENVNAAMKAASNESYGYTEDPI  
VSSDIVGMSYGSFLDATQTKVLDVDGKQLVKVVSWDNEMSYTAQLVRTLGLLRKNC

**SP036 nucleotide (SEQ ID NO:55)**

TTCTTACGAGTTGGGACTGTATCAAGCTAGAACGGTTAAGGAAAATAATCGTGTTCCTATATAGATGG  
AAAACAAGCGACGCAAAAAACGGAGAATTTGACTCCTGATGAGGTTAGCAAGCGTGAAGGAATCAATGC  
TGAGCAAATCGTCATCAAGATAACAGACCAAGGCTATGTCACTTCACATGGCGACCCTATCATTATTA  
CAATGGTAAGGTTCTTATGACGCTATCATCAGTGAAGAATTACTCATGAAAGATCCAACTATAAGCT  
AAAAGATGAGGATATTGTTAATGAGGTCAAGGGTGGATATGTTATCAAGGTAGATGGAAAACTATGT  
TTACCTTAAGGATGCTGCCACGCGGATAACGTCCGTACAAAAGAGGAAATCAATCGACAAAAACAAGA  
GCATAGTCAACATCGTGAAGGTGGAACCTCAAGAAACGATGGTGTCTTGCCTTGGCACGTTTCGAAGG  
ACGCTATATCTACAGATGATGGTTATATCTTTAATGCTTCTGATATCATAGAGGATACTGGTGATGCTTA  
TATCGTTCCTCATGGAGATCATTACCATTACATCTCTTAAGAATGAGTTATCAGCTAGCGAGTTGGCTGC  
TGCAGAAGCCTTCCTATCTGGTTCGAGGAAATCTGTCAAATTCAAGAACCCTATCGCCGACAAAATAGCGA  
TAACACTTCAAGAACAACTGGGTACCTTCTGTAAGCAATCCAGGAACCTACAAATACTAACACAAGCAA  
CAACAGCAACACTAACAGTCAAGCAAGTCAAAGTAATGACATTGATAGTCTCTTGAAACAGCTCTACAA  
ACTGCCCTTGAGTCAACGACATGTAGAATCTGATGGCCTTGTCTTTGATCCAGCACAAATCACAAGTCG  
AACAGCTAGAGGTGTTGCAGTGCCACACGGAGATCATTACCCTTCATCCCTTACTCTCAAATGTCTGA  
ATTGGAAGAACGAATCGCTCGTATTATTTCCCTTCGTTATCGTTCAAACCATTGGGTACCAGATTCAAG  
GCCAGAACCAAGTCCACAACCGACTCCGGAACCTAGTCCAGGCCCGCAACCTGCACCAAATCTTAA  
AATAGACTCAAATCTCTCTTTGGTTAGTCAGCTGGTACGAAAAGTTGGGGAAGGATATGTATTGGAAGA  
AAAGGGCATCTCTCGTTATGTCTTTGCGAAAGATTTACCATCTGAAACTGTTAAAAATCTTGAAAGCAA  
GTTATCAAAAACAAGAGAGTGTTCACACACTTTAACTGCTAAAAAAGAAAATGTTGCTCCTCGTGACCA  
AGAATTTTATGATAAAGCATATAATCTGTTAACTGAGGCTCATAAAGCCTTGTTTGNAAAATAAGGGTCG  
TAATCTGATTTCGAAGCCTTAGACAAATTATTAGAAGCCTTGAATGATGAATCGACTAATAAAGAAAA  
ATTGGTAGATGATTATTGGCATTCTTAGCACCATTACCCATCCAGAGCGACTTGGCAAACCAAATTC  
TCAAATTGAGTATACTGAAGACGAAGTTCGTATTGCTCAATTAGCTGATAAGTATACAACGTCAGATGG  
TTACATTTTGTATGAACATGATATAATCAGTGATGAAGGAGATGCATATGTAACGCCTCATATGGGQCA  
TAGTCACTGGATTGGAAAAGATAGCCTTTCTGATAAGGAAAAAGTTGCAGCTCAAGCCTATACTAAAGA  
AAAAGGTATCCTACCTCCATCTCCAGACGCAGATGTTAAAGCAAATCCAACTGGAGATAGTGCAGCAGC  
TATTTACAATCGTGTGAAAGGGGAAAAACGAATTCACCTCGTTCGACTTCCATATATGGTTGAGCATAC  
AGTTGAGGTTAAAAACGGTAATTTGATTATTCCTCATAAGGATCATTACCATAATATTAAATTTGCTTG  
GTTTGATGATCACACATACAAAGCTCCAAATGGCTATACCTTGGAAGATTTGTTTGCGACGATTAAGTA  
CTACGTAGAACACCCTGACGAACGTCCACATCTAATGATGGATGGGGCAATGCCAGTGAGCATGTGTT  
AGGCAAGAAAGACCACAGTGAAGATCCAAATAAGAACTTCAAAGCGGATGAAGAGCCAGTAGAGGAAAC  
ACCTGCTGAGCCAGAAGTCCCTCAAGTAGAGACTGAAAAAGTAGAAGCCCAACTCAAAGAAGCAGAAGT  
TTTGCTTGCGAAAGTAACGGATTCTAGTCTGAAAGCCAAATGCAACAGAACTCTAGCTGGTTTACGAAA  
TAATTTGACTCTTCAAATTATGGATAACAATAGTATCATGGCAGAAGCAGAAAAATTACTTGCGTTGTT  
AAAAGGAAGTAATCCTTCATCTGTAAGTAAGGAAAAAATAAAC

**SP036 amino acid (SEQ ID NO:56)**

SYELGLYQARTVKENNRVSYIDGKQATQKTENLTPDEVSKREGINAEQIVIKITDQGYVTSHGDHYHY  
NGKVPYDAIISEELMKDPNYKLKDEDIVNEVKGGYVIKVDGKYVYLKDAHADNVRTKEEINRQKQE

Table 1

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HSQHREGGTPRNDGAVALARSQGRYTTDDGYIFNASDIIEDTGDAYIVPHGDHYHYIPKNELSASELAA  
AEAFLSGRGNLSNSRTYRRQNSDNTSRTNWVPSVSNPGTTNTNTSNNSTNSQASQSNDDIDSLKQLYK  
LPLSQRHVESDGLVFDPAQITSRTARGVAVPHGDHYHFIPYSQMSELEERIIPLRYRSNHWPDSR  
PEQSPQPTEPSPGPQPAPNLKIDSNSSLVSQLVRKVGEYVFEKGISRYVFAKDLPSETVKNLESK  
LSKQESVSHTLTAKKENVAPRDQEFYDKAYNLLTEAHKALFXNKGRNSDFQALDKLLERLNDESTNKEK  
LVDDLLAFLAPITHPERLGKPNISQIEYTEDEVRIAQLADKYTTSDGYIFDEHDIISDEGDAYVTPHMGH  
SHWIGKDSLSDKEKVAAQAYTKEKGILPPSPDADVKANPTGDSAAAIYNRVKGEKRIPLVRLPYMVEHT  
VEVKNGNLIIPH KDHYHNIKFAWFDHITYKAPNGYTLEDLFATIKYYVEHPDERPHSNDGWGNASEHVL  
GKKDHSDEPNKNFKADEEPVEETPAEPEVPQVETEKVEAQLKEAEVLLAKVTDSSLKANATETLAGLRN  
NLTLQIMDNNSIMAEAEKLLALLKGSNPSSVSKEKIN

**SP038 nucleotide (SEQ ID NO:57)**

TACTGAGATGCATCATAATCTAGGAGCTGAAAAGCGTTCAGCAGTGGCTACTACTATCGATAGTTTTAA  
GGAGCGAAGTCAAAAAGTCAGAGCACTATCTGATCCAAATGTGCGTTTTGTTCCTTCTTTGGCTCTAG  
TGAATGGCTTCGTTTTGACGGTGCTCATTCTGCGGTATTAGCTGAGAAATACAATCGTTCCTACCGTCC  
TTATCTTTTAGGACAGGGGGAGCTGCATCGCTTAACCAATATTTTGGAAATGCAACAGATGTTACCACA  
GCTGGAGAATAAACAAGTTGTGTATGTTATCTCACCTCAGTGGTTCAGTAAAAATGGCTATGATCCAGC  
AGCCTTCCAGCAGTATTTTAATGGAGACCAAGTTGACTAGTTTTCTGAAACATCAATCTGGGGATCAGGC  
TAGTCAATATGCAGCGACTCGCTTACTGCAACAGTTCCTCAAACGTAGCTATGAAGGACCTGGTTCAGAA  
GTTGGCAAGTAAAGAAGAATTGTGACAGCAGACAATGAAATGATTGAATATTGGCTCGTTTTTAATGA  
ACGCCAAGCTTCCTTTTTTGGTCAGTTTTTCGGTTAGAGGCTATGTTAACTACGATAAGCATGTAGCTAA  
GTATTTAAAAATCTTGCCAGACCAGTTTTCTTATCAGGCAATAGAAGATGTTGTCAAAGCAGATGCTGA  
AAAAATACTTCCAATAATGAGATGGGAATGGAAAATTATTTCTATAATGAGCAGATCAAGAAGGATTT  
GAAGAAATTAAGGATTCTCAGAAAAGCTTTACCTATCTCAAGTCGCCAGAGTATAATGNNTTGCAGTT  
GGTTTTAACACAGTTTTCTAAATCTAAGGTAAACCCGATTTTTTATCATTCCACCTGTTAATAAAAAATG  
GATGNACTATGCTGGTCTACGAGAGGATATGTACCAACAAACGGTGCAGAAGATTCGCTACCAGTTAGA  
AAGTCAAGGTTTTTACCAATATAGCAGATTTTTCTAAGGACGGCGGGGAGCCTTTCTTTATGAAGGACAC  
CATTCACTTGGTTGGTTGGTTGGCTTTTGACAAGGCAGTTGATCCTTTCTCTATCCAATCCCAC  
ACCAGCTCCGACTTACCATCTGAATGAGCGCTTTTTTCAGCAAAGATTGGGCGACTTATGATGGAGATGT  
CAAAGAA

**SP038 amino acid (SEQ ID NO:58)**

TEMHHNLGAEKRSVAVATTIDSFKERSQKVRALSDPNVRFVPPFGSSEWLRFDAHSAVLAEKYNSYRYP  
YLLGQGAASLNQYFGMQMLPQLENKQVVYVISPQWFSKNGYDPAAFQQYFNGDQLTSFLKHQSGDQA  
SQYAATRLQQFPNVAMKDLVQKLASKEELSTADNEMIELLARFNERQASFFGQFSVRGVVNYDKHVAK  
YLKILPDQFSYQAIEDVVKADEKNTSNNEMGMENYFYNEQIKKDLKKLSDSKSFTYLLKSPYENXLQL  
VLTQFSKSKVNPFIPIPPVNKKWMXYAGLREDMYQQTQVQKIRYQLESQGFTNIADFSKDGGEPPFMKDT  
IHLGWLGWLAFDKAVDPFLSNPTPPTYHLNERFFSKDWATYDGDVKE

**SP039 nucleotide (SEQ ID NO:59)**

GGTTTTGAGAAAGTATTTGCAGGGGGCCCTGATTGAGTCGATTGAGCAAGTGGAAGTACCGTATTGT  
GGAAATTACAGTTTCCAATAAAAACGAGATTGGAGACCATATCCAGGCTACCTTGATTATCGAAATTAT  
GGGGAAACACAGTAATATTCTACTGGTCGATAAAAGCAGTCATAAAATCCTCGAAGTTATCAAACACGT  
CGGCTTTTCACAAAATAGCTACCGCACCTTACTTCCAGGATCGACCTATATCGCTCCGCCAAGTACAAA  
ATCTCTCAATCCTTTTACTACTAAGGATGAAAAGCTCTTTGAAATCCTGCAAACCAAGAACTAACAGC  
AAAAATCTTCAAAGCCTCTTTCAAGGTCTGGGACGCGATACGGCAAATGAATTGGAAAGGATACTGGT  
TAGTGAAAAACTTTCCGCTTTCCGAAATTTTTTCAATCAAGAAACCAAGCCATGCTTGACTGAGACTTC  
CTTCAGTCCAGTTCTTTTGCAAATCAGGTGGGAGAGCCTTTTGCAAATCTTTCTGATTTGTTGGACAC  
CTACTATAAGGATAAGGCTGAGCGCGACCGCGTCAAACAGCAGGCCAGTGAAGTATTCGTCGTGTTGA  
AAATGAAGTTCAGAAAAACCGACACAACTCAAAAAACAGGAAAAAGAGTTACTGGCGACAGACAACGC  
TGAAGAATTCGTCAAAAAGGAGAATTGCTGACAACCTTCTCCACCAAGTGCCTAACGACCAAGACCA  
GGTTATCTAGACAATACTATAACCAACCAACCTATCATGATTGCGCTTGATAAGGCTCTGACTCCCAA  
CCAGAATGCCCAACGCTATTTTAAACGGTATCAGAACTCAAAGAAGCTGTCAAATACTTGACTGATTT  
GATTGAAGAAACCAAGCCACTATTCTCTATCTGGAAAGTGTAGAAACCGTCTCAACCAAGCTGGACT  
GGAAGAAATCGCTGAAATCCGTGAAGAATTGATTCAAACAGGTTTTTATCCGCAGAAGACAACGGGAGAA  
AATCCAGAAACGCAAAAACTAGAACAAATATCTAGCAAGCGATGGCAAAACCATCATCTATGTCGGACG  
AAACAATCTTCAAATGAGGAATTGACCTTTAAATGGCCCGCAAGGAGGAAGTTTGGTTCCATGCTAA  
GGACATTCCTGGAAGCCATGTTGTCTATCTCAGGAAATCTTGACCCATCTGATGCAGTCAAGACAGACGC



Table 1

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AGCAGAGTTAGCTGCCTACTTCTCTCAAGGGCGCCTGTGCAATCTGGTGCAGGTAGATATGATTGAAGT  
CAAAAACTCAATAAACCACTGGTGGAAAACCCGGCTTTGTCACTTACACAGGACAAAAGACCCCTCCG  
CGTCACACCAGACTCCAAAAAATTGCATCCATGAAAAATCC

**SP039 amino acid (SEQ ID NO:60)**

VLRKYLQGALIESIEQVENDRIVEITVSNKNEIGDHIQATLIIIEIMGKHSNILLVDKSSHKILEVIKHY  
GFSQNSYRTLLPGSTYIAPPSTKSLNPFTIKDEKLFEILQTQELTAKNLQSLFQGLGRDTANELERILV  
SEKLSAFRNFFNQETKPCLTETSFSPVPFANQVGEFFANLSDLLDTYYKDKAERDRVKQQAASELIRVE  
NELQKNRHLKKQEKELLATDNAEEFRQKGELLTTFHLHQVPNDQDQVILDNYYTNQPIIMIALDKALTPN  
QNAQRYFKRYQKLKEAVKYLTDLIEETKATILYLESVETVLNQAGLEEIAEIREELIQTGFIRRRQREK  
IQKRKKLEQYLASDGKTIIVYGRNNLQNEELTFKMARKEELWFHAKDIPGSHVVISGNLDPDAVKTD  
AELAAFYFSGRLSNLVQVDMIEVKKLNKPTGGKPGFVITYTGQKTLRVTPDSKKIASMKKS

**SP040 nucleotide (SEQ ID NO:61)**

GACAACATTTACTATCCATACAGTAGAGTCAGCACCAGCAGAGAAGTGAAAGAAATTCCTGAAACAGTAGA  
AAAAGACAACAATGGCTATATTCCCAACCTAATCGGTCTCTTGGCCAATGCCCGACTGTTTTAGAACG  
CTACCAAATTGTCTCATCTATCCACCGTCGCAACAGCCTGACACCCGTTGAGCGTGAAGTGGTGCAAAAT  
CACGGCAGCCGTGACCAATGGTTGTGCCTTCTGTGTGCGAGGTACACAGCCTTTTCCATCAAAACAAAT  
CCAGATGAATGATGACTTGATTCAAGCTCTTCGCAATCGTACTCCAATTGAAACAGATCCTAAATTGGA  
TACCCTAGCTAAGTTTACCTTGGCAGTTATCAATACCAAGGGTCGTGTAGGAGATGAAGCCTTGTCTGA  
GTTTTTAGAAGCTGGCTACACTCAACAAAATGCCTTGGATGTGGTTTTTGGTGTGAGCCTAGCAATCCT  
CTGTAATATGCCAACAACTTAGCTAATACCAATTAATCCAGAATTGCAACCTTATGCC

**SP040 amino acid (SEQ ID NO:62)**

TTFTIHTVESAPAEVKEILETVEKDNNGYIPNLIGLLANAPTVEAYQIVSSIHRNSLTPVEREVVQI  
TAAVTNGCAFCVAGHTAFSIKQIQMNDLILQALRNRTPIETDPKLDLAKFTLAVINTKGRVGDALSE  
FLEAGYTQQNALDVVFGVSLAILCNYANNLANTPINPELQPYA

**SP041 nucleotide (SEQ ID NO:63)**

GGCTAAGGAAAGAGTGATGTACTAGCTTATAAACAGGGGTTGTTTGAAACGAGAGAGCAGGCCAAGCG  
AGGTGTGATGGCTGGCCTAGTCGTAGCAGTCCTTAATGGAGAACGGTTTGACAAGCCAGGAGAGAAAAAT  
TCCAGATGACACCGAATTAATACTCAAGGGGGAGAACTCAAGTATGTCAGCCGTGGTGGTTTTGAACT  
GGAAAAGGCCTTGCAAGTCTTTGATTTGTGCGGTGGATGGCGCGACTACGATTGATATCGGGGCCTCTAC  
TGGAGGTTTTACCGATGTCATGCTACAGAATAGTGCCAAGTTGGTCTTTGCAGTCGATGTTGGTACCAA  
TCAGTTGGCTTGAAATTACGCCAAGACCCACGAGTTGTCAGCATGGAGCAGTTCAATTTCCGCTATGC  
TGAAAAGACTGATTTTCGAGCAGGAGCCGAGCTTTGCCAGTATTGATGTGAGTTTCATTCCCTTAGTCT  
GATTTTGGCAGCCTTGACCCGTGTCTTGGCTGATCAAGGTCAGGTGGTAGCACTTGTCAAACCTCAGTT  
TGAGGCAGGACGTGAGCAGATTGGGAAAAATGGAATTATTCGAGATGCTAAGGTTTCATCAGAATGTCCT  
TGAATCTGTAACAGCTATGGCAGTAGAGGTAGGTTTTTCAGTCCTTGGCTTGGACTTTTCTCCCATCCA  
AGGTGGACATGGAAATATTGAATTTTTCAGCTATTTGAAAAAGAAAAGTCAGCAAGCAATCAGATTCT  
TGCTGAGATTAAAGAAGCAGTAGAGAGGGCGCATAGTCAATTTAAAAATGAA

**SP041 amino acid (SEQ ID NO:64)**

AKERVDVLAYKQGLFETREQAKRGVMAGLVVAVLNGERFDKPGEKIPDDTELKLGKELKYVSRGGLKL  
EKALQVFDLSVDGATTIDIGASTGGFTDVMLQNSAKLVFAVDVGTNQLAWKLQDPRVVSMEQFNFRYA  
EKTDFEQEPSFASIDVSFISLSLILPALHRVLADQGGVVALVKPQFEAGREQIGKNGIIRDAKVHQNVL  
ESVTAMAVEVGFVSLGLDFSPIQGGHGNIEFLAYLKKEKSASNQILAEIKEAVERAHSQFKNE

**SP042 nucleotide (SEQ ID NO:65)**

TTGTTCCATGAACCTGGTCTGCACCAAGCTGGTCAGGTTAAGAAAGAGTCTAATCGAGTTTCTTATAT  
AGATGGTGATCAGGCTGGTCAAAAGGCAGAAAACCTTGACACCAGATGAAGTCAGTAAGAGGGAGGGGAT  
CAACGCCGAACAAATNGTNATCAAGATTACGGATCAAGGTTATGTGACCTCTCATGGAGACCATTATCA  
TTACTATAATGGCAAGGTTTCTTATGATGCCATCATCAGTGAAGAGCTCCTCATGAAAGATCCGAATTA  
TCAGTTGAAGGATTGACACATTGTCAATGAAATCAAGGGTGGTTATGTGATTAAAGGTAAACGGTAAATA  
CTATGTTNACCTTAAGGATGCAGCTCATGCGGATAATATTCGGACAAAAGAAGAGATTAAACGTCAGAA  
GCAGGAACGCAGTCATAATCATAACTCAAGAGCAGATAATGCTGTTGCTGCAGCCAGAGCCCAAGGACG  
TTATACAACGGATGATGGGTATATCTTCAATGCATCTGATATCATTGAGGACACGGGTGATGCTTATAT  
CGTTCCTCACGGCGACCATTACCATTACATTCCTAAGAATGAGTTATCAGCTAGCGAGTTAGCTGCTGC

Table 1

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AGAAGCCTATTGGAATGGGAAGCAGGGATCTCGTCCTTCTTCAAGTTCTAGTTATAATGCAAATCCAGC  
TCAACCAAGATTGTGAGAGAACCACAATCTGACTGTCACTCCAACTTATCATCAAAATCAAGGGGAAAA  
CATTTCAAGCCTTTTACGTGAATTGTATGCTAAACCTTATCAGAACGCCATGTGGAATCTGATGGCCT  
TATTTTCGACCCAGCGCAAATCACAAGTCGAACCGCCAGAGGTGTAGCTGTCCCTCATGGTAACCATTA  
CCACTTTATCCCTTATGAACAAATGTCTGAATTGGAAAAACGAATTGCTCGTATTATTCCTTCGTTA  
TCGTTCAAACCATTTGGGTACCAGATTCAAGACCAGAACAACCAAGTCCACAATCGACTCCGGAACCTAG  
TCCAAGTCCGCAACCTGCACCAATCCTCAACCAGCTCCAAGCAATCCAATTGATGAGAAATTGGTCAA  
AGAAGCTGTTTCGAAAAGTAGGCGATGGTTATGTCTTTGAGGAGAATGGAGTTTCTCGTTATATCCCAGC  
CAAGGATCTTTTCAGCAGAAACAGCAGCAGGCATTGATAGCAAACTGGCCAAGCAGGAAAGTTTATCTCA  
TAAGCTAGGAGCTAAGAAAACAGCTCCCATCTAGTATCGAGAATTTTACAATAAGGCTTATGACTT  
ACTAGCAAGAATTACCAAGATTTACTTGATAATAAAGGTCGACAAGTTGATTTTGAGGCTTTGGATAA  
CCTGTTGGAACGACTCAAGGATGTNCNAAGTGATAAAGTCAAGTTAGTGGANGATATTCTTGCCTTCTT  
AGCTCCGATTTCGTATCCAGAAGCTTTAGGAAAAACCAATGCGCAAATTACCTACACTGATGATGAGAT  
TCAAGTAGCCAAGTTGGCAGGCAAGTACACAACAGAAGACGGTTATATCTTTGATCCTCGTGATATAAC  
CAGTGATGAGGGGGATGCCTATGTAACCTCCACATATGACCCATAGCCACTGGATTAAAAAAGATAGTTT  
GTCTGAAGCTGAGAGAGCGGCAGCCAGGCTTATGCTAAAGAGAAAGGTTTGACCCCTCCTTCGACAGA  
CCATCAGGATTTCAGGAAATACTGAGGCAAAAGGAGCAGAAGCTATCTACAACCGCGTGAAAGCAGCTAA  
GAAGGTGCCACTTGATCGTATGCCTTACAATCTTCAATATACTGTAGAAGTCAAAAACGGTAGTTTAAT  
CATACCTCATTATGACCATTACCATACATCAAAATTTGAGTGGTTTGACGAAGGCCCTTTATGAGGCACC  
TAAGGGGTATACCTTTGAGGATCTTTTGCGGACTGTCAAGTACTATGTGCAACATCCAAACGAACGTC  
GCATTTCAGATAATGGTTTGGTAACGCTAGCGACCATGTTCAAAGAAACAAAAATGGTCAAGCTGATAC  
CAATCAAACGGAAAAACCAAGCGAGGAGAAACCTCAGACAGAAAAACCTGAGGAAGAAACCCCTCGAGA  
AGAGAAACCGCAAAGCGAGAAACAGAGTCTCCAAAACCAACAGAGGAACCAGAAGAATCACCAGAGGA  
ATCAGAAGAACCTCAGGTCGAGACTGAAAAGGTTGAAGAAAACTGAGAGAGGCTGAAGATTTACTTGG  
AAAAATCCAGGAT

**SP042 amino acid (SEQ ID NO:66)**

CSYELGRHQAGQVKESNRVSYIDGDQAGQKAENLTPEVSKREGINAEQXVIKITDQGYVTSBGDHYH  
YYNGKVPYDAIIESELLMKDPNYQLKDSDIVNEIKGGYVIKVNKYYVYLKDAHADNIRTKEEIKRQK  
QERSHNHNSRADNAVAARAQGRYTTDDGYIFNASDIIEDTGDAYIVPHGDHYHYI PKNELSASELAAA  
EAYWNGKQGSRPSSSSSYNANPAQPRLSENHNLTVTPYHQNGENISSLLRELYAKPLSERHVESDGL  
IFDPAQITSRTARGVAVPHGNHYHFIPEYQMSLEKRIARIIPLYRSNHWVPDSRPEQPSQPSTPEPS  
PSPQPAPNPQPAPSNPIDELVKEAVRKVG DG YVFEENGVSRYIPAKDLSAETAAGIDSKLAKQESLSH  
KLGAKKTDLPSSDREFYNKAYDLLARIHQDLLDNKGRQVDFEALDNLLERLKD VXS D K V K L V X D I L A F L  
APIRHPERLGPNAQITYTDDIEIQVAKLAGKYTTEDGYIFDPRDITSDEGDAYVTPHMTSHWIKKDSL  
SEAERAAAQAYAKEKGLTPPSTDHQSNGTEAKGAEAIYNRVKA AKK VPLDRMPYNLQYTV EVKNGSLI  
IPHYDHYHNIKFEWFDGLYEAPKGYTLEDLLATVKYVVEHPNERPHSDNGFGNASDHVQRNKNQADT  
NQTEKPSEPKPQTEKPEEETPREPKPQSEKPESPKPTEPEESPEESEEPQVETEKVEEKLREAEDLLG  
KIQD

**SP043 nucleotide (SEQ ID NO:67)**

TTATAAGGGTGAATTAGAAAAAGGATACCAATTTGATGGTTGGGAAATTTCTGGTTTCGAAGGTAAAAA  
AGACGCTGGCTATGTTATTAATCTATCAAAAGATACCTTTATAAAACCTGTATTCAAGAAAAATAGAGGA  
GAAAAAGGAGGAAGAAAAATAAACCTACTTTTGATGTATCGAAAAAGAAAGATAACCCACAAGTAAACCA  
TAGTCAATTAAATGAAAGTCACAGAAAAGAGGATTTACAAAGAGAAGAGCATTACAAAAATCTGATTCT  
AACTAAGGATGTTACAGCTACAGTTCTTGATAAAAACAATATCAGTAGTAAATCACTACTAACAATCC  
TAATAAG

**SP043 amino acid (SEQ ID NO:68)**

YKGELEKGYQFDGWEISGFEGKKDAGYVINLSKDTFIKPVFKKIEEKKEEENKPTFDVSKKKDNPQVNH  
SQLNESHKEDLQREEHSQKSDSTKDV TATVLDKNNISSKSTTNNPNK

**SP044 nucleotide (SEQ ID NO:69)**

GAATGTTTCAGGCTCAAGAAAGTTTCAGGAAATAAAATCCACTTTATCAATGTTCAAGAAGGTGGCAGTGA  
TGCGATTATTCTTGAAAGCAATGGACATTTTGCCATGGTGGATACAGGAGAAGATTATGATTTCCAGGA  
TGGAAGTGATTCTCGCTATCCATGGAGAGAAGGAATTGAAACGTCTTATAAGCATGTTCTAACAGACCG  
TGCTTTTCGTCGTTTGAAGGAATTGGGTGTCCAAAACCTTGATTTTATTTTGGTGACCCATACCCACAG  
TGATCATATTGAAATGTTGATGAATTACTGTCTACCTATCCAGTTGACCGAGTCTATCTTAAGAAATA

Table 1

TAGTGATAGTCGTATTACTAATTCTGAACGTCTATGGGATAATCTGTATGGCTATGATAAGGTTTTACA  
GACTGCTGCAGAAAAAGGTGTTTCAGTTATTCAAATATCACACAAGGGGATGCTCATTTCAGTTTGG  
GGACATGGATATTACAGCTCTATAATTATGAAAATGAACTGATTCATCGGGTGAATTAAAGAAAATTTG  
GGATGACAATTCCAATTCCTTGATTAGCGTGGTGAAAGTCAATGGCAAGAAAATTTACCTTGGGGGCGA  
TTTAGATAATGTTTCATGGAGCAGAAGACAAGTATGGTCCTCTCATTGGAAAAGTTGATTTGATGAAGTT  
TAATCATCACCATGATACCAACAAATCAAATACCAAGGATTTCAATAAAAATTTGAGTCCGAGTTTGAT  
TGTTCAAACCTTCGGATAGTCTACCTTGAAAAATGGTGTGATAGTGAGTATGTTAATTGGCTCAAAGA  
ACGAGGAATTGAGAGAATCAACGCAGCCAGCAAAGACTATGATGCAACAGTTTTTGATATTCGAAAAGA  
CGGTTTTGTCAATATTTCAACATCCTACAAGCCGATCCAAGTTTTCAAGCTGGTTGGCATAAGAGTGC  
ATATGGGAACCTGGTGGTATCAAGCGCTGATTCTACAGGAGAGTATGCTGTGCGTTGGAATGAAATCGA  
AGGTGAATGGTATTACTTTAAACCAAACGGGTATCTTGTTACAGAATCAATGGAAAAATGGAACAATCA  
TTGGTTCTATTGACAGACTCTGGTGCTTCTGCTAAAAATTTGGAAGAAAATCGCTGGAATCTGGTATTA  
TTTTAACAAAGAAAACAGATGGAAATTGGTTGGATTCAAGATAAAGAGCAGTGGTATTATTTGGATGT  
TGATGGTTCTATGAAGACAGGATGGCTTCAATATATGGGGCAATGGTATTACTTTGCTCCATCAGGGGA  
A

**SP044 amino acid (SEQ ID NO:70)**

NVQAQESSGNKIHFINVQEGGSDAIILESNGHFAMVDITGEDYDFPDGSDSRYPWREGIETSYKHVLTDR  
VFRRLKELGVQKLDLILVTHSDHIGNVDELLSTYPVDRVYLKKYSDSRITNSERLWDNLGYDKVLQ  
TAAEKGVSVIQNITQGDHAFQFGDMDIQLYNYENETDSSGELKKIWDDNSNSLISVVKVNGKKIYLGGD  
LDNVHGAEDKYGPLIGKVDLMKFNNHHDNKSNTKDFIKNLSPSLIVQTSDSL PWKNGVDSEYVNWKE  
RGIERINAASKDYDATVFDIRKDG FVNISTSYKPIPSFQAGWHKSAYGNWWYQAPDSTGEYAVGWNEIE  
GEWYYFNQTGILLQNWKKWNHWFYLTDSGASAKNWKKIAGIWIYYFNKENQMEIGWIQDKEQWYYLDV  
DGSMTGWLQYMGQWYFAPSGE

**SP045 nucleotide (SEQ ID NO:71)**

CTTGGGTGTAACCCATATCCAGCTCCTTCCAGTCTTGTCTTACTACTTTGTCAATGAATTGAAAAACCA  
TGAACGCTTGTCTGACTACGCTTCAAGCAACAGCAACTACAACCTGGGGATATGACCTCAAACTACTT  
CTCCTTGACTGGTATGTACTCAAGCGATCCTAAGAATCCAGAAAAACGAATCGCAGAATTTAAACCT  
CATCAACGAAATCCACAAACGTGGTATGGGAGCTATCCTAGATGTCGTTTATAACCACACAGCCAAAGT  
CGATCTCTTTGAAGATTTGGAACCAAACCTACTACCCTTTATGGATGCCGATGGCACACCTCGAAGTAG  
CTTTGGTGGTGGACGCTTGGGGACAACCCACCATATGACCAAACGGCTCCTAATTGACTCTATCAAATA  
CCTAGTTGATACCTACAAAGTGGATGGCTTCCGTTTCGATATGATGGGAGACCATGACGCCGCTTCTAT  
CGAAGAAGCTTACAAGGCTGCACGCGCCCTCAATCCAAACCTCATCATGCTTGGTGAAGGTTGGAGAAC  
CTATGCCGCTGATGAAAACATGCCTACTAAAGCTGCTGACCAAGATTGGATGAAACATACCGATACTGT  
CGCTGTCTTTTCAGATGACATCCGTAACAACCTCAAATCTGGTTATCCAAACGAAGGTCAACCTGCCTT  
TATCACAGGTGGCAAGCGTGATGTCAACACCATCTTTAAAAATCTCATTGCTCAACCAACTAAGTTTGA  
AGCTGACAGCCCTGGAGATGTCATCCAATACATCGCAGCCCATGATAACTTGACCTCTTTGACATCAT  
TGCCAGTCTATCAAAAAGACCCAAGCAAGGCTGAGAATATGCTGAAATCCACCGTCGTTTACGACT  
TGGAATCTCATGGTCTTGACAGCTCAAGGAATCCATTTATCCACTCCGGTCAGGAATATGGACGTAC  
TAAACAATTCGCTGACCCAGCCTACAAGACTCCAGTAGCAGAGGATAAGGTTCCAAACAAATCTCACTT  
GTTGCGTGATAAGGACGGCAACCCATTGACTATCCTTACTTCATCCATGACTCTTACGATTCTAGTGA  
TGCAGTCAACAAGTTTGACTGGACTAAGGCTACAGATGGTAAAGCTTATCCTGAAAATGTCAAGAGCCG  
TGACTATATGAAAGGTTTGATTGCCCCCTTCGTCAATCTACAGATGCCTTCCGACTTAAGAGTCTTCAAGA  
TATCAAAGACCGTGTCACCTCATCACTGTCCAGGCCAAAAATGGTGTGGAAGAGGATGTAGTGAT  
TGGCTACCAAATCACTGCTCCAAACGGCGATATCTACGCAGTCTTTGTCAATGCGGATGAAAAGCTCG  
CGAATTTAATTTGGGAAGTGCCTTTGCACATCTAAGAAATGCGGAAGTTTTGGCAGATGAAAACCAAGC  
AGGACAGTCGGAATTGCCAACCCGAAAGGACTTGAATGGACTGAAAAAGGCTTGAAATTGAATGCCCT  
TACAGCTACTGTTCTTCGAGTCTCTCAAATGGAAGTACCATGAGTCAACTGCAGAAGAGAAACCAGA  
CTCAACCCCTTCCAAGCCTGAACATCAAATGAAGCTTCTCACCTGCACATCAAGACCCAGCTCCAGA  
AGCTAGACCTGATTCTACTAAACCAGATGCCAAAGTAGCTGATGCGGAAAATAAACCTAGCCAAGCTAC  
AGCTGATTCAAGCTGAACAACCAGCACAGAAGCACAAAGCATCATCTGTAAAAGAAGCGGTTTCGAAA  
CGAATCGGTAGAAAACCTTAGCAAGGAAAATATACCTGCAACCCCAAGATAAACAAGCTGAA

**SP045 nucleotide (SEQ ID NO:72)**

LGVTHIQLLPVLSTYFVNELKNHERLSDYASSNSNYNWGYDPQNYFSLTGMYSDDPKNPEKRIAEFKNL  
INEIHKRGMGAILDVVYNHTAKVDLFEDLEPNYYHFMDADGTPRTSFGGGRLGTTHHMTKRLLIDSIFY  
LVDYTKVDGFRFDDMMGDHDAASIEEAYKAARALNPNLIMLGEGWRTYAGDENMPTKAADQDWMKHTDTV

Table 1

AVFSDDIRNNLKSYPNEGQPAFITGGKRDVNTIFKNLIAQPTNFEADSPGDVIQYIAAHNLTFLFDII  
AQSIKKDPSKAENYAEIHRRLRLGNLMVLTAQGTPFIHSGQEYGRTKQFRDPAYKTPVAEDKVPNKSHL  
LRDKDGNPFDPYFIHDSYDSSDAVNKFDTWKATDGKAYPENVKSRDYMKGILALRQSTDAFRLKSLQD  
IKDRVHLITVPGQNGVEKEDVIGYQITAPNGDIYAVFVNADEKAREFNLGTAFALRNAEVLADENQA  
GPVGIANPKGLEWTEKGLKLNALTATVLRVSQNGTSHESTAEKPDSTPSKPEHQNEASHPAHQDPAPE  
ARPDSTKPDPAKVADAENKPSQATADSQAEQPAQEAQASSVKEAVRNESVENSSKENIPATPDKQAE

**SP046 nucleotide (SEQ ID NO:73)**

TAGTGATGGTACTTGGCAAGGAAAAAGTATCTGAAAGAAGATGGCAGTCAAGCAGCAAATGAGTGGGT  
TTTNGATACTCATTATCAATCTTGGTTCTATATAAAAGCAGATGCTAACTATGCTGAAAATGAATGGCT  
AAAGCAAGGTGACGACTATTTTTACCTCAAATCTGGTGGCTATATGGCCAAATCAGAATGGGTAGAAGA  
CAAGGGAGCCTTTTATTATCTTGACCAAGATGGAAGATGAAAAGAAATGCTTGGGTAGGAACTTCCTA  
TGTTGGTGCAACAGGTGCCAAAAGTAATAGAAGACTGGGTCTATGATTCTCAATACGATGCTTGGTTTAA  
TATCAAAGCAGATGGACAGCAGCAGAGAAAAGTGGCTCCAAATTAAAGGGAAGGACTATTATTTCAA  
ATCCGGTGGTTATCTACTGACAAGTCAGTGGATTAATCAAGCTTATGTGAATGCTAGTGGTGCCAAAAGT  
ACAGCAAGGTTGGCTTTTTGACAAAACAATACCAATCTTGGTTTTACATCAAAGAAAATGGAAACTATGC  
TGATAAAGAATGGATTTTCGAGAATGGTCACTATTATTATCTAAAATCCGGTGGCTACATGGCAGCCAA  
TGAATGGATTTGGGATAAGGAATCTTGGTTTTATCTCAAATTTGATGGGAAAATGGCTGAAAAAGAATG  
GGTCTACGATTTCTCATAGTCAAGCTTGGTACTACTTCAAATCCGGTGGTTACATGACAGCCAAATGAATG  
GATTTGGGATAAGGAATCTTGGTTTTACCTCAAATCTGATGGGAAAATAGCTGAAAAAGAATGGGTCTA  
CGATTTCTCATAGTCAAGCTTGGTACTACTTCAAATCTGGTGGCTACATGGCGAAAATGAGACAGTAGA  
TGGTTATCAGCTTGGAAGCGATGGTAAATGGCTTGGAGGAAAAACTACAAATGAAAATGCTGCTTACTA  
TCAAGTAGTGCTGTACAGCCAAATGTTTATGATTAGATGGTGAAGGCTTTCCTATATATCGCAAGG  
TAGTGTCTGATGGCTAGATAAGGATAGAAAAAGTGATGACAAGCGCTAGATGCTAGTAAGGACTTTATCCCTTATTATGA  
GAGTGATGGCCACCGTTTTTATCATTATGTGGCTCAGAATGCTAGTATCCCAGTAGCTTCTCATCTTTC  
TGATATGGAAGTAGGCAAGAAATATTATTCCGCAGATGGCCTGCATTTTGATGGTTTTAAGCTTGAGAA  
TCCCTTCTTTTTCAAAGATTTAACAGAGGCTACAAACTACAGTGTGAAGAATTGGATAAGGTATTTAG  
TTTGCTAAACATTAACAATAGCCTTTTGGAGAACAAGGGCGCTACTTTTAAGGAAGCCGAAGAACATTA  
CCATATCAATGCTCTTTATCTCCTTGCCCATAGTGCCCTAGAAAGTAAGTGGGGAAGAAAGTAAATTC  
CAAAGATAAGAATAATTTCTTTGGCATTACAGCCTATGATACGACCCCTTACCTTTCTGCTAAGACATT  
TGATGATGTGATAAGGGAATTTTAGGTGCAACCAAGTGGATTAAGGAAAATTATATCGATAGGGGAAG  
AACTTTCTTGGAACAAGGCTTCTGGTATGAATGTGGAATATGCTTCAGACCCTTATTGGGGCGAAAA  
AATTGCTAGTGTGATGATGAAAATCAATGAGAAGCTAGGTGGCAAAGAT

**SP046 amino acid (SEQ ID NO:74)**

SDGTWQGKQYLKEDGSQAANEVXDTHYQSWFYIKADANYAENEWLKQDDYFYLKSGGYMAKSEWVED  
KGAFYYLDQDGKMKRNAWVGTSYVGATGAKVIEDWVYDSQYDAWFIKADGQHAKEWLQIKGKDYFYK  
SGGYLLTSQWINQAYVNASGAKVQGWLFQKQYQSWFYIKENGNADKEWIFENGHYLYLKSGGYMAAN  
EWIWDKESWFIKFDGKMAEKWVYDSHSQAWYFYKSGGYMTANEWIWDKESWFIKSDGKIAEKWVY  
DSHSQAWYFYKSGGYMAKNETVDGYQLGSDGKWLGGKTTNENAAYYQVVPVTANVYDSGDEKLSYISQG  
SVVWLDKDRKSDDKRLAITISGLSGYMKTEDLQALDASKDFIPYYESDGHFRFYHYVAQNASIPVASHLS  
DMEVGKKYYSADGLHFDGFKLENPFLFKDLTEATNYSAEELDKVFSLLNINNSLLENKGATFKEAEEHY  
HINALYLLAHSALSNWGRSKIADKNNFFGITAYDTPYLSAKTFDDVDKGILGATKWIKENYIDRGR  
TFLGNKASGMNVEYASDPYWGEKIASVMMKINEKLGGKD

**SP048 nucleotide (SEQ ID NO:75)**

TGGGATTCAATATGTCAGAGATGATACTAGAGATAAAGAAGAGGGAATAGAGTATGATGACGCTGACAA  
TGGGGATATTATTGTAAAAGTAGCGACTAAACCTAAGGTAGTAACCAAGAAAATTTCAAGTACGCGAAT  
TCGTTATGAAAAAGATGAAACAAAAGACCGTAGTGAAAATCCTGTTACAATTGATGGAGAGGATGGCTA  
TGTAACACGACAAGGACCTACGATGTTAATCCAGAGACTGGTTATGTTACCGAACAGGTTACTGTTGA  
TAGAAAAGAAGCCACGGATACAGTTATCAAAGTCCAGCTAAAAGCAAGGTTGAAGAAGTTCTTGTTC  
ATTTGCTACTAAATATGAAGCAGACAATGACCTTTCTGCAGGACAGGAGCAAGAGATTACTCTAGGAAA  
GAATGGGAAAACAGTTACAACGATAACTTATAATGTAGATGGAAAGAGTGGACAAGTAAGTGAAGTAC  
TTTAAGTCAAAAAAAGACTCTCAAACAAGAGTTGTTAAAAAAGAACCAkCCCCAAGTTCTTGTCCA  
AGAAATTCCAATCGAAACAGAATATCTCGATGGCCCACTCTTGATAAAAAGTCAAGAAGTAGAAGAAGT  
AGGAGAAATTGGTAAATTACTCTTACTACAATCTATACTGGTAGATGAACGTGATGGAACAATTGAAGA  
AACTACTTCTCGTCAAATTACTAAAGAGATGGTAAAAAGACGTATAAGGAGAGGGACGAGAGAACCTGA

Table 1

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AAAAGTTGTTGTTCCCTGAGCAATCATCTATTCCTTCGTATCCTGTATCTGTTACATCTAACCAAGGAAC  
AGATGTAGCAGTAGAACCAGCTAAAGCAGTTGCTCCAACAACAGACTGGAAACAAGAAAATGGTATGTG  
GTATTTTTATAATACTGATGGTTCCATGGCAACAGGTTGGGTACAAGTTAATAGTTCATGGTACTACCT  
CAACAGCAACGGTTCTATGAAAGTCAATCAATGGTTCCAAGTTGGTGGTAAATGGTATTATGTAAATAC  
ATCGGGTGAGTTAGCGGTCAATACAAGTATAGATGGCTATAGAGTCAATGATAATGGTGAATGGGTGCC  
T

**SP048 amino acid (SEQ ID NO:76)**

GIQYVRDDTRDKEEGIEYDDADNGDIIVKVATKPKVVTKKISSSTRIRYEKDETKDRSENPVITIDGEDGY  
VTTTRTYDVNPETGYVTEQVTVDRKEATDTVIKVPKSKVEEVLVPFATKYEADNDLSAGQEIEITLGK  
NGKTVTTITYNVGKSGQVTESTLSQKKDSQTRVVKKRTXPQVLVQEIPIETEYLDGPTLSDKSQEVEEV  
GEIGKLLLLLSILVDERDGTIEETTSRQITKEMVKRRIRRGRTREPEKVVVPEQSSIPSYPSVTSNQGT  
DVAVEPAKAVAPTDDWKQENGMWYFYNTDGSMATGWVQVNSSWYYLNSNGSMKVNQWFQVGGKWYVNT  
SGELAVNTSIDGYRVNDNGEVR

**SP049 nucleotide (SEQ ID NO:77)**

GGATAATAGAGAAGCATTAAAAACCTTTATGACGGGTGAAAATTTTTATCTCCAACATTATCTAGGAGC  
ACATAGGGAAGAACTAAATGGAGAGCATGGCTATACCTTCCGTGTTTGGGCACCTAATGCTCAGGCTGT  
TCACTTGGTTGGTGATTTTACCAACTGGATTGAAAATCAGATTCCAATGGTAAGAAATGATTTTGGGGT  
CTGGGAAGTCTTTACCAATATGGCTCAAGAAGGGCATATTTACAAATATCATGTCACACGTCAAAATGG  
TCATCAACTGATGAAGATTGACCCTTTGTCTGTCAGGTATGAGGCTCGTCCAGGAACAGGGGCAATCGT  
AACAGAGCTTCTGAGAAGAAATGGAAGGATGGACTTTGGCTGGCAGCAAGAAAACGTTGGGGCTTTGA  
AGAGCGTCTGTCAATATTTATGAAGTTCACGCTGGATCATGGAAGAAATTTCTGATGGCAGTCCTTA  
TAGTTTTGCCAGCTCAAGGATGAAGTCAATTCCTTATCTCGTTGAAATGAAGTATACTCATATTGAGTT  
TATGCCCTTGATGTCCCATCCTTTGGGCTTGAGTTGGGGGTATCAGCTTATGGGTTACTTCGCTTTAGA  
GCATGCTTATGGCCGACCAGAGGAGTTTCAAGATTTTGTCT

**SP049 amino acid (SEQ ID NO:78)**

DNREALKTFMTGENFYLQHYLGAHREELNGEHGYTFRVWAPNAQAVHLVGDFTNWIENQIPMVRNDFGV  
WEVFTNMAQEGHIYKYHVTRQNGHQLMKIDPFAVRYEARPGTGAIIVTELPEKKWKDGLWLARRKRWFGE  
ERPVNIYEVHAGSWKRNSDGSPYSFAQLKDELIPYLVEMNYTHIEFMPLMSHPLGLSWGYYQLMGYFALE  
HAYGRPEEFQDFV

**SP050 nucleotide (SEQ ID NO:79)**

AGATTTTGTGCGAGGAGTGTACATACCCATAATATTGGGGTTATTGTGGACTGGGTACCAGNTCACTTTAC  
CATCAACGATGATGCCTTAGCCTATTATGATGGGACACCGACTTTTGAATACCAAGACCATAATAAGGC  
TCATAACCATGGTTGGGGTGCCCTTAATTTTGACCTTGGAAGAAATGAAGTCCAGTCCCTTCTTAATTTT  
TTGCATTAAGCATTGGATTGATGTCTATCATTTGGATGGTATTCTGTGGATGCTGTTAGCAACATGCT  
CTATTGGACTATGATGATGCTCCATGGACACCTAATAAAGATGGCGGAAATCTCAACTATGAAGGTTA  
TTATTTCCCTTCAGCGCTTGAATGAGGTTATTAAGTTAGAATATCCAGATGTGATGATGATTGCAGAAGA  
AAGTTTCGTCTGCGATCAAGATTACGGGAATGAAAGAGATTGGTGGTCTAGGATTTGACTACAAATGGAA  
CATGGGCTGGATGAATGATATCCTCCGTTTCTACGAAGAAGATCCGATCTATCGTAAATATGACTTTAA  
CCTGGTGACTTTTCACTTTATGATATGTTTCAAGGAGAATTATCTCTTGCCATTCTCGCAGCATGAAGT  
GGTTCATGGCAAGAAGAGTATGATGCATAAGATGTGGGGAGATCGTTACAATCAATTCGCAGGCTTGCG  
CAATCTCTATACGTACCAATTTGTCAACCTGGTAAGAAATTGCTCTTCATGGGTAGCGAATACGGTCA  
ATTCCTAGAATGGAATCTGAAGAACAGTTGGAATGGTCTAACCTAGAAGACCCAATGAATGCTAAGAT  
GAAGTATTTTCGCTTCTCAGCTAAACAGTTTACAAAGATCATCGCTGTCTGTGGGAAATTGATACCAG  
CTATGATGGTATTGAAATCATTGATGCGGATAATCGAGACCAGAGTGTTCTTTCTTTTATTTCGTAAGGG  
TAAAAAGGGA

**SP050 amino acid (SEQ ID NO:80)**

DFVEECHTHNIGVIVDWVPXHFTINDDALAYYDGTPTFEYQDHNKAHNHGWGALNFDLGKNEVQSFLIS  
CIKHWDIVYHLDGIRVDAVSNNMLYLDYDDAPWTPNKDGGNLYEGYYFLQRLNEVIKLEYPDVMMAEE  
SSSAIKITGMKEIGGLGFDYKWNMGWMNDILRFYEDPIYRKYDFNLVTF SFMYVXKENYLLPFSHDEV  
VHGKKSMMHKMWGDRYNQFAGLRNLYTYQICHGPKLLFMGSEYQGFLEWKSEEQLEWSNLEDPMNAKM  
KYFASQLNQFYKDHRLWEIDTSYDGIIEIIDADNRDQSVLSFIRKGKKG

**SP051 nucleotide (SEQ ID NO:81)**

Table 1

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ATCTGTAGTTTATGCGGATGAAACACTTATTACTCATACTGCTGAGAAACCTAAAGAGGAAAAAATGAT  
 AGTAGAAGAAAAGGCTGATAAAGCTTTGGAACTAAAAATATAGTTGAAAGGACAGAACAAAGTGAACC  
 TAGTTCAACTGAGGCTATTGCATCTGAGNAGAAAAGAAGATGAAGCCGTAACCTCCAAAAGAGGAAAAAGT  
 GTCTGCTAAACCGGAAGAAAAAGCTCCAAGGATAGAATCACAAGCTTCAAATCAAGAAAAACCGCTCAA  
 GGAAGATGCTAAAGCTGTAACAAATGAAGAAGTGAATCAAATGATTGAAGACAGGAAAGTGGATTTTAA  
 TCAAAATTTGGTACTTTAAACTCAATGCAAAATCTAAGGAAGCCATTAAACCTGATGCAGACGTATCTAC  
 GTGGAAAAAATTAGATTTACCGTATGACTGGAGTATCTTTAACGATTTTCGATCATGAATCTCCTGCACA  
 AAATGAAGGTGGACAGCTCAACGGTGGGGAAGCTTTGGTATCGCAAGACTTTTCAAACCTAGATGAAAAAGA  
 CCTCAAGAAAAATGTTTCGCCCTTACTTTTGTATGGCGTCTACATGGATTCTCAAGTTTATGTCAATGGTCA  
 GTTAGTGGGGCATTATCCAAATGGTTATAACCAGTTCTCATATGATATCACCAAATACCTTCAAAAAGA  
 TGGTCGTGAGAATGTGATTGCTGTCCATGCAGTCAACAAACAGCCAAGTAGCCGTTGGTATTACAGGAAG  
 TGGTATCTATCGTGATGTGACTTTACAAGTGACAGATAAGGTGCATGTTGAGAAAAATGGGACAACCTAT  
 TTTAACACCAAAACTTGAAGAACAACAACATGGCAAGGTTGAAACTCATGTGACCAGCAAAATCGTCAA  
 TACGGACGACAAAGACCATGAACCTGTAGCCGAATATCAAATCGTTGACGAGGTGGTGCATGCTGTAA  
 AGGCTTAGTTTCGTACAGCGAGTTCGTACCTTAAAAAGCACATGAATCAACAAGCCCTAGATGCGATTTTGA  
 AGTTGAAAGACCAAACTCTGGACTGTTTTAAATGACAAACCTGCCTTGTACGAATTGATTACGCGTGT  
 TTACCGTGACGGTCAATTGGTTGATGCTAAGAAGGATTTGTTTGGTTACCGTTACTATCACTGGACTCC  
 AAATGAAGGTTTCTCTTTGAATGGTGAACGTATTAAATTCATGGAGTATCCTTGCACCACGACCATGG  
 GGCGCTTGGAGCAGAAGAAAACATAAAGCAGAATATCGCCGCTCTCAAACAAATGAAGGAGATGGGAGT  
 TAACTCCATCCGTACAACCCACAACCTGCTAGTGAGCAAACCTTGCAAATCGCAGCAGAACTAGGTTT  
 ACTCGTTCAGGAAGAGGCCCTTTGATACGTGGTATGGTGGCAAGAAACCTTATGACTATGGACGTTTCTT  
 TGA AAAAGATGCCACTCACCCAGAAGCTCGAAAAGGTGAAAAATGGTCTGATTTTGACCTACGTACCAT  
 GGTCCGAAAGAGGCAAAAACAACCTGCTATCTTCATGTGGTCAATTGGTAATGAAATAGGTGAAGCTAA  
 TGGTGATGCCCCTCTTTAGCAACTGTTAAACGTTTGGTTAAGGTTATCAAGGATGTTGATAAGACTCG  
 CTATGTTACCATGGGAGCAGATAAATTCCTGTTTCGGTAATGGTAGCGGAGGGCATGAGAAAATTGCTGA  
 TGAACCTCGATGCTGTTGGATTAACTATTTCTGAAGATAATTACAAAGCCCTTAGAGCTAAGCATCCAAA  
 ATGGTTGATTTATGGATCAGAAACATCTTCAGCTACCCGTACACGTGGAAGTTACTATCGCCCTGAACG  
 TGAATTGAAACATAGCAATGGACCTGAGCGTAATTATGAACAGTCAGATTATGGAAATGATCGTGTGGG  
 TTGGGGGAAAACAGCAACCGCTTCATGGACTTTTGACCGTGACAACGCTGGCTATGCTGGACAGTTTAT  
 CTGGACAGGTACGGACTATATTGGTGAACCTACACCATGGCACAACCAAAATCAAACCTCCTGTTAAGAG  
 CTCTTACTTTGGTATCGTAGATACAGCCGGCATTCCAAAACATGACTTCTATCTCTACCAAAGC

**SP051 amino acid (SEQ ID NO:82)**

SVVYADETLITHAEKPKKEKMIVEEKADKALETKNIVERTEQSEPSSTEAIASEXKEDEAVTPKEEKV  
 SAKPEEKAPRIESQASNQEKPLKEDAKAVTNEEVNQMIEDRKVDFNQNWYFKLNANSKEAIKPDADVST  
 WKKLDLPYDWSIFNDFDHESPAQNEGGQLNGGEAWYRKTFKLDEKDLKKNVRLTFDGVYMDSQVYVNGQ  
 LVGHYPNGYNQFSYDITKYLQKDGRENVIAVHAVNKPSSRWYSGSGIYRDVTLQVTDKVVHVEKNGTTI  
 LTPKLEEQQHKGKVETHVTSKIVNTDDKDHELVAEYQIVERGGHAVTGLVRTASRTLKAHESTSLDAILE  
 VERPKLWTVLNDKPALYELITRVYRDGQLVDAKKDLFGYRYHWTNPNEGFSLNGERIKFHGVSLHHDHG  
 ALGAEENYKAEYRRLKQMKEMGVNSIRTHNPASEQTLQIAELGLLVQEEAFDTWYGGKKPYDYGRFF  
 EKDATHPEARKEKWSDFDLRTMVERGKNPFIWMFSIGNEIGEANGDAHSLATVKRLVKVIKDVKTR  
 YVTMGADKFRFGNGSGGHEKIADELDAVGFNYSYEDNYKALRAKHPKWLIIYGETSSATRTRGSYYRPER  
 ELKHSNGPERNYEQSDYGNDRVGWGTATASWTFDRDNAGYAGQFIWTGTDYIGEPTPWHNQNTPVKS  
 SYFGIVDTAGIPKHDFYLYQS

**SP052 nucleotide (SEQ ID NO:83)**

TTACTTTGGTATCGTAGATACAGCCGGCATTCCAAAACATGACTTCTATCTCTACCAAAGCCAATGGGT  
 TTCTGTAAAGAAGAAACCGATGGTACACCTTCTTCTCACTGGAAGTGGGAAAACAAAGAATTAGCATC  
 CAAAGTAGCTGACTCAGAAGGTAAGATTCCAGTTCGTGCTTATTTCGAATGCTTCTAGTGTAGAATTGTT  
 CTTGAATGGAAAATCTCTTGGTCTTAAAGACTTTCAATAAAAAACAACAGCGATGGGCGGACTTACCA  
 AGAAGGTGCAAATGCTAATGAACCTTTATCTTGAATGGAAAGTTGCCATCAACCAGGTACCTTGGAGC  
 AATTGCTCGTGATGAATCTGGCAAGGAAATGCTCGAGATAAGATTACGACTGCTGGTAAGCCAGCGGC  
 AGTTCGCTCTATTAAAGGAAGACCATGCGATTCGAGCAGATGGAAAAGACTTGACTTACATCTACTATGA  
 AATTGTTGACAGCCAGGGGAATGTGGTTCCAACCTGCTAATAATCTGGTTTCGCTTCCAATTGCATGGCCA  
 AGGTCAACTGGTCCGTGTAGATAACGGAGAACAAGCCAGCCGTGAACGCTATAAGGCGCAAGCAGATGG  
 TTCTTGGATTTCGTAAAGCATTTAATGGTAAAGGTGTTGCCATTGTCAAATCAACTGAACAAGCAGGGAA  
 ATTCACCCTGACTGCCCACTCTGATCTCTTGAATCGAACCAAGTCACTGTCTTTACTGGTAAGAAAGA  
 AGGACAAGAGAAGACTGTTTTGGGGACAGAAGTGCCAAAAGTACAGACCATTATTGGAGAGGCACCTGA

Table 1

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AATGCCTACCACTGTTCCGTTTGTATACAGTGATGGTAGCCGTGCAGAACGTCCTGTAACCTGGTCTTC  
AGTAGATGTGAGCAAGCCTGGTATTGTAACGGTGAAAGGTATGGCTGACGGACGAGAAGTAGAAGCTCG  
TGTAAGAAGTGATTGCTCTTAAATCAGAGCTACCAAGTTGTGAAACGTATTGCTCCAAATACTGACTTGAA  
TTCTGTAGACAAATCTGTTTCCATGTTTGTGATTGATGGAAGTGTGAAAGAGTATGAAGTGACAAAGTG  
GGAGATTGCCGAAGAAGATAAAGCTAAGTTAGCAATTCCAGGTTCTCGTATTCAAGCGACCGGTTATTT  
AGAAGGTCAACCAATTTCATGCAACCCCTTGTGGTAGAAGAAGGCAATCCTGCGGCACCTGCAGTACCAAC  
TGTAACGGTTGGTGGTGAGGCAGTAACAGGTCTTACTAGTCAAAAACCAATGCAATACCGCACTCTTGC  
TTATGGAGCTAAGTTGCCAGAAGTCACAGCAAGTGCTAAAAATGCAGCTGTACAGTTCTTCAAGCAAG  
CGCAGCAAACGGCATGCGTGCGAGCATCTTTATTACGCTAAAGATGGTGGCCCTCTTCAAACCTATGC  
AATTCAATTCTTTGAAGAAGCGCCAAAAATTGCTCACTTGAGCTTGCAAGTGGAAGCTGACAGTCT  
CAAAGAAGACCAAACCTGTCAAATTGTCGGTTCGAGCTCACTATCAAGATGGAACGCAAGCTGTATTACC  
AGCTGATAAAGTAACCTTCTCTACAAGTGGTGAAGGGGAAGTCGCAATTCTGTAAGGAATGCTTGAGTT  
GCATAAGCCAGGAGCAGTCACTCTGAACGCTGAATATGAGGGAGCTAAAGACCAAGTTGAACCTACTAT  
CCAAGCCAATACTGAGAAGAAGATTGCGCAATCCATCCGTCCTGTAAATGTAGTGACAGATTTGCATCA  
GGAACCAAGTCTTCCAGCAACAGTAACAGTTGAGTATGACAAAGGTTTCCCTAAAACTCATAAAGTCAC  
TTGGCAAGCTATTCCGAAAGAAAACTAGACTCCTATCAAACATTTGAAGTACTAGGTAAAGTTGAAGG  
AATTGACCTTGAAGCGCGTGCAAAAGTCTCTGTAGAAGGTATCGTTTTCAGTTGAAGAAGTCAGTGTGAC  
AACTCCAATCGCAGAAGCACCACAATTACCAGAAAGTGTTCGGACATATGATTCAAATGGTCACGTTTC  
ATCAGCTAAGGTTGCATGGGATGCGATTCTGCCAGAGCAATACGCTAAGGAAGGTGTCTTTACAGTTAA  
TGGTCGCTTAGAAGGTACGCAATTAACA

**SP052 amino acid (SEQ ID NO:84)**

YFGIVDTAGIPKHDFFLYQSQWVSVKKKPMVHLLPHWNWENKELASKVADSEGKIPVRAYSNNASSVELF  
LNGKSLGLKTFNKKQTS DGRTYQEGANANELYLEWKVAYQPGTLEAIARDESGKEIARDKITTAGKPAA  
VRLIKEDHAIADGKDLTYIYYEIVDSQGNVVPANLVRFLHGGQQLVGVDNGEQASRERYKAQADG  
SWIRKAFNGKGVAIKSTEQAGKFTLTAHSDLLKSNQVTVFTGKKEGQEKTVLGTVEVPKVQTIIGEAPE  
MPTTVPFVYSDGSRAERPVTWSSVDVSKPGIIVTVKGMADGREVEARVEVIALKSELPVVKRIAPNTDLN  
SVDKSVSYVLIDGSVEEYEVDKWEIAEEDKAKLAIPGSRIQATGYLEGQPIHATLVVEEGNPAAPAVPT  
VTVGSEAVTGLTSLQKPMQYRTLAYGAKLPEVTASAKNAAVTVLQASAANGMRASIFIQPKDGGPLQTYA  
IQFLEEAPKIAHLSLQVEKADSLKEDQTVKLSVRAHYQDGTQAVLPADKVTFTS SGEGEVAIRKGMLEL  
HKPGAVTLNAEYEGAKDQVELTIQANTEKKIAQSIRPVNVVTDLHQEPSLPATVTVEYDKGFPKTHKVT  
WQAIPEKLDLSYQTFEVLGKVEGIDLEARAKVSVEGIVSVEEVSVTTPIAEAPQLPESVRTYDSNGHVS  
SAKVANDAIRPEQYAKEGVFTVNGRLEGTQLT

**SP053 nucleotide (SEQ ID NO:85)**

AGCTAAGGTTGCATGGGATGCGATTCTGCCAGAGCAATACGCTAAGGAAGGTGTCTTTACAGTTAATGG  
TCGCTTAGAAGGTACGCAATTAACAACCTAACTTTCATGTTTCGCGTATCTGCTCAAACCTGAGCAAGGTGC  
AAACATTTCTGACCAATGGACCGGTTTCAGAAATTGCCACTTGCCCTTTGCTTCAGACTCAAATCCAAGCGA  
CCCAGTTTCAAATGTTAATGACAAGCTCATTTCTACAATAACCAACCAGCCAATCGTTGGACAAACTG  
GAATCGTACTAATCCAGAAGCTTCAGTCGGTGTCTGTTTGGAGATTACAGGTATCTTGAGCAAACGCTC  
CGTTGATAATCTAAGTGTGCGATTCCATGAAGACCATGGAGTTGGTGTACCGAAGTCTTATGTGATTGA  
GTATTATGTTGGTAAGACTGTCCCAACAGCTCCTAAAAACCTAGTTTTGTTGGTAATGAGGACCATGT  
CTTTAATGATTCTGCCAAGTGGAAACAGTTACTAATCTAAAAGCCCCCTGCTCAACTCAAGGCTGGAGA  
AATGAACCACTTTAGCTTTTGATAAAGTTGAAACCTATGCTGTTTCGTATTTCGCATGGTTAAAGCAGATAA  
CAAGCGTGGAACGTCTATCACAGAGGTACAAATCTTTGCGAAACAAGTTGCGGCAGCCAAGCAAGGACA  
AACAAGAATCCAAGTTGACGGCAAAGACTTAGCAAACTTCAACCCTGATTTGACAGACTACTACCTTGA  
GTCTGTAGATGGAAAAGTTCCGGCAGTCACAGCAAGTGTAGCAACAATGGTCTCGCTACCGTCGTTCC  
AAGCGTTCGTGAAGGTGAGCCAGTTCGTGTATCGCGAAAGCTGAAAATGGCGACATCTTAGGAGAATA  
CCGCTCTGCACTTCACTAAGGATAAGAGCTTACTTTCTCATAAACCAGTTGCTGCGGTTAAACAAGCTCG  
CTTGCTACAAGTAGGTCAAGCACTTGAATTGCCGACTAAGGTTCCAGTTTACTTCACAGGTAAGACGG  
CTACGAAACAAAAGACCTGACAGTTGAATGGGAAGAAGTTCCAGCGGAAAATCTGACAAAAGCAGGTCA  
ATTTACTGTTCGAGGCCGTGTCTTGGTAGTAACCTTGTGCTGAGATCACTGTACGAGTGACAGACAA  
ACTTGGTGAGACTCTTTTCAGATAACCCTAACTATGATGAAAACAGTAACCAGGCCTTTGCTTCAGCAAC  
CAATGATATTGACAAAACTCTCATGACCGCGTTGACTATCTCAATGACGGAGATCATTGACAAAATCG  
TCGTTGGACAAACTGGTCACCAACACCATCTTCTAATCCAGAAGTATCAGCGGTTGTGATTTTCCGTGA  
AAATGGTAAGATTGTAGAACGGACTGTTACACAAGGAAAAGTTTCAGTTCTTTGCAGATAGTGGTACGGA  
TGCACCATCTAAACTCGTTTTAGAACGCTATGTGCGTCCAGAGTTTGAAGTGCCAACCTACTATTCAA  
CTACCAAGCCTACGACGCAGACCATTCAACAATCCAGAAAATTTGGGAAGCTGTTCCTTATCGTGC

Table 1

GGATAAAAGACATTGCAGCTGGTGATGAAATCAACGTAACATTTAAAGCTATCAAAGCCAAAGCTATGAG  
ATGGCGTATGGAGCGTAAAGCAGATAAGAGCGGTGTTGCGATGATTGAGATGACCTTCCTTGACCAAG  
TGAATTGCCTCAAGAAAGCACTCAATCAAAGATTCTTGTTAGATGGAAAAGAACTTGCTGATTTGCTGA  
AAATCGTCAAGACTATCAAATTACCTATAAAGGTCAACGGCCAAAAGTCTCAGTTGAAGAAAACAATCA  
AGTAGCTTCAACTGTGGTAGATAGTGGAGAAGATAGCTTTCCAGTACTTGTTCGCCTCGTTTCAGAAAG  
TGGAAAACAAGTCAAGGAATACCGTATCCACTTGACTAAGGAAAAACCAGTTTCTGAGAAGACAGTTGC  
TGCTGTACAAGAAGATCTTCCAAAATCGAATTTGTTGAAAAAGATTTGGCATACAAGACAGTTGAGAA  
AAAAGATTCAACACTGTATCTAGGTGAAACTCGTGTAGAACAAGAAGGAAAAGTTGGAAAAGAACGTAT  
CTTTACAGCGATTAATCCTGATGGAAGTAAGGAAGAAAACTCCGTGAAGTGGTAGAAGTTCCGACAGA  
CCGCATCGTCTTGGTTGGAACCAACCAGTAGCTCAAGAAGCTAAAAAACCACAAGTGTGAGAAAAAGC  
AGATACAAAACCAATTGATTCAAGTGAAGCTAGTCAAACTAATAAAGCCCAG

**SP053 amino acid (SEQ ID NO:86)**

AKVAWDAIRPEQYAKEGVFTVNGRLEGTQLTTLKLVHVRVSAQTEQGANISDQWTGSELPLAFASDSNP  
PVSNNVDKLISYNNQPANRWTNWNRTNPEASVGVLFGDSGILSKRSVDNLSVGFHEDHGVGVPKSYVIE  
YYVGKTVPTAPKNPSFVGNEDHVFNDSANWKPVNTLKAQAQLKAGEMNHF SFDKVETAVRIRMVKADN  
KRGTSITEVQIFAKQVAAAKQGQTRIQVDGKDLANFNPDLDYYLESVGDKVPAVTASVSNNGLATVVP  
SVREGEPPVRVIAKAENGDI LGEYRLHFTKDKSLLSHKPVAAVKQARLLQVQALELPTKVPVYFTGKDG  
YETKDLTVÉWEEVPAENLT KAGQFTVRGRVLGSLNVAEITVRVTDKLGELSDNPNYDENSNAFASAT  
NDIDKNSHDRVLDYLDGDHSENRRWTNWSPTPSSNPEVSAGVIFRENGKIVERTVTQGVQFFADSGTD  
APSKLVLERYVGPFEFVPTYYSNYQAYDADHPFNNPENWEAVPYRADKDIAAGDEINVTFKAIKAKAMR  
WRMERKADKSGVAMIEMTFLAPSEL PQESTQSKILVDGKELADFAENRQDYQITYKGQRPKVSVEENNQ  
VASTVVDSDGDSFPVLVRLVSESGKVKEYRIHLTKEKPVSEKTVAAVQEDLPKIEFVEKDLAYKTVEK  
KDSTLYLGETRVEQEGKVGERIFTAINPDGSKEEKLREVVEVPTDRIVLVGTPVAQEAKPKVQVSEKA  
DTKPIDSSEASQTNKAQ

**SP054 nucleotide (SEQ ID NO:87)**

CTATCACTATGTAAATAAAGAGATTATTTCAACAAGAAGCTAAAGATTTAATTCAGACAGGAAAGCCTGA  
CAGGAATGAAGTTGTATATGGTTTGGTGTATCAAAAAGATCAGTTGCCTCAAACAGGGACAGAA

**SP054 amino acid (SEQ ID NO:88)**

YHYVNKEIISQEAKDLIQTKPDRNEVVYGLVYQKDQLPQTGTE

**SP055 nucleotide (SEQ ID NO:89)**

TGAGACTCCTCAATCAATAACAAATCAGGAGCAAGCTAGGACAGAAAACCAAGTAGTAGAGACAGAGGA  
AGCTCCAAAAGAAGAAGCACCTAAAACAGAAGAAAGTCCAAAGGAAGAACC AAAATCGGAGGTAAAACC  
TACTGACGACACCTTCCTAAAGTAGAAGAGGGGAAAGAAGATTCAGCAGAACCAAGCTCCAGTTGAAGA  
AGTAGGTGGAGAAGTTGAGTCAAAACCAGAGGAAAAGTAGCAGTTAAGCCAGAAAAGTCAACCATCAGA  
CAAACCAGCTGAGGAATCAAAAGTTGAACAAGCAGGTGAACCAGTCGCGCCAAGAGAAGACGAAAAGGC  
ACCAGTCGAGCCAGAAAAGCAACCAGAAGCTCCTGAAGAAGAGAAGGCTGTAGAGGAAACACCGAAACA  
AGAAGAGTCAACTCCAGATACCAAGGCTGAAGAACTGTAGAACC AAAAGAGGAGACTGTTAATCAATC  
TATTGAACAACCAAAAGTTGAAACGCCTGCTGTAGAAAAACAAACAGAACCAACAGAGGAACCAAAAGT  
TGAACAAGCAGGTGAACCAGTCGCGCCAAGAGAAGACGAACAGGCACCAACGGCACCAGTTGAGCCAGA  
AAAGCAACCAGAAGTTCTTGAAGAAGAGAAGGCTGTAGAGGAAACACCGAAACCAGAAGATAAAATAAA  
GGGTATTGGTACTAAAGAACCAGTTGATAAAAGTGAGTTAAATAATCAAATTGATAAAGCTAGTTTCAGT  
TTCTCCTACTGATTAT

**SP055 amino acid (SEQ ID NO:90)**

ETPQSITNQEQARTENQVVETEAPKEEAPKTEESPKEEPSEVKPTDDTLPKVEEGKEDSAEPAPVEE  
VGGEVESKPEEKVAVKPESQPSDKPAEESKVEQAGEPVAPREDEKAPVEPEKQPEAPEEEKAVEETPKQ  
EESTPDTKAEETVEPKETVNQSI EQPKVETPAVEKQTEPTEEPKVEQAGEPVAPREDEQAPTAPVEPE  
KQPEVP EEKAVEETPKPEDKIKGIGTKEPVDKSELNNQIDKASSVSPTDY

**SP056 nucleotide (SEQ ID NO:91)**

GGATGCTCAAGAAACTGCGGGAGTTCACTATAAATATGTGGCAGATTGAGAGCTATCATCAGAAGAAAA  
GAAGCAGCTTGTCTATGATATTCGACATACGTGGAGAAATGATGATGAACTTATTATCTTGTATTATAA  
GTTAAATTCTCAAATCAACTGGCGGAATTGCCAAATACTGGAAGCAAGAATGAGAGGCAA



Table 1

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## SP056 amino acid (SEQ ID NO:92)

DAQETAGVHYKYVADSELSSEEKKQLVYDIPTYVENDDETYYLVYKLNSQNQLAELPNTGSKNERQ

## SP057 nucleotide (SEQ ID NO:93)

CGACAAAGGTGAGACTGAGGTTCAACCAGAGTCGCCAGATACTGTGGTAAGTGATAAAGGTGAACCAGA  
GCAGGTAGCACCCTTCCAGAAATATAAGGGTAATATTGAGCAAGTAAAACCTGAAACTCCGGTTGAGAA  
GACCAAAGAACAAGGTCCAGAAAAAACTGAAGAAGTTCCAGTAAAACCAACAGAAGAAACACCAGTAAA  
TCCAAATGAAGGTACTACAGAAGGAACCTCAATTCAAGAAGCAGAAAAATCCAGTTCAACCTGCAGAAGA  
ATCAACAACGAATTACAGAGAAAAGTATCACCAGATACATCTAGCAAAAATACTGGGGAAGTGTCCAGTAA  
TCCTAGTGATTTCGACAACCTCAGTTGGAGAATCAAATAAACAGAACATAATGACTCTAAAAATGAAAA  
TTCAGAAAAAACTGTAGAAGAAAGTTCCAGTAAATCCAAATGAAGGCACAGTAGAAGGTACCTCAAATCA  
AGAAACAGAAAAACCAGTTCAACCTGCAGAAGAAACACAAACAACTCTGGGAAAATAGCTAACGAAAA  
TACTGGAGAAGTATCCAATAAACCTAGTGATTCAAACACCAGTTGAAGAATCAAATCAACCAGAAAA  
AAACGGAAGTGAACAAAAACCAGAAAAATTCAGGTAATACAACATCAGAGAATGGACAAACAGAACCCAGA  
ACCATCAAACGGAAATTCAACTGAGGATGTTTCAACCGAATCAAACACATCCAATTCAAATGGAACCGA  
AGAAATTAAACAAGAAAAATGAACTAGACCCTGATAAAAAGGTAGAAGAACCAGAGAAAAACACTTGAATT  
AAGAAAT

## SP057 amino acid (SEQ ID NO:94)

DKGETEVQPESPDTVVSDKGEPEQVAPLPEYKGNIEQVKPETPVEKTKEQGPEKTEEVVKPTEETPVN  
PNEGTTGEGTSIQEENPVQPAEESTTNSEKVPDTSKNTGEVSSNPDSSTTSVGESNKPEHNDKSKNEN  
SEKTVEEVPVNPNEGTVGEGTSNQETEKPVQPAEETQTN SGKIANENTGEVSNKPSDKPPVEESNQPEK  
NGTATKPENSGNTTSENGQTEPEPSNGNSTEDVSTESNTSNSNGNEEIKQENELDPDKKVEEPEKTLER  
RN

## SP058 nucleotide (SEQ ID NO:95)

AAATCAATTGGTAGCACAAAGATCCAAAAGCACAAAGATAGCACTAAACTGACTGCTGAAAAATCAACTGT  
TAAAGCACCCTGCTCAAAGAGTAGATGTAAAAGATATAACTCATTAAACAGATGAAGAAAAAGTTAAGGT  
TGCTATTTTACAAGCAAATGGTTTCAGCATTAGACGGAGCGACAATCAATGTAGCTGGAGATGGTACAGC  
AACAATCACATTCCAGATGGTTTCAGTAGTGACGATTCTAGGAAAAGATACAGTTCAACAATCTGCGAA  
AGGTGAATCTGTAAGTCAAGAAGCTACACCAGAGTATAAGCTAGAAAAATACACCAGGTGGAGATAGGG  
AGGCAATACTGGAAGCTCAGATGCTAATGCGAATGAAGGCGGTGGTAGCCAGGCGGGTGGATCAGCTCA  
CACAGGTTCAAAAACCTCAGCTCAATCACAAGCTTCTAAGCAATTAGCTACTGAAAAAGAATCAGCTAA  
AAATGCCATTGAAAAAGCAGCCAAGGACAAGCAGGATGAAATCAAAGGCGCACCGCTTTCTGATAAAGA  
AAAAGCAGAACTTTTAGCAAGAGTGGAAGCAGAAAAACAAGCAGCTCTCAAAGAGATTGAAATGCGAA  
AACTATGGAAGATGTGAAGGAAGCAGAAACGATTGGAGTGCAAGCCATTGCCATGGTTACAGTTCCTAA  
GAGACCACTGGCTCCTAAT

## SP058 amino acid (SEQ ID NO:96)

NQLVAQDPKAQDSTKLTAEKSTVKAPAQRVDVKDITHLTDEEKVKVAILQANGSALDGATINVAGDGTA  
TITFPDGSVVTILGKDTVQQSAKGESVTQEATPEYKLENTPGGDKGGNTGSSDANANEGGGSQAGGSAH  
TGSQNSAQSQASKQLATEKESAKNAIEKAAKDKQDEIKGAPLSDKEKAELLARVEAEKQAALKEIENAK  
TMEDVKEAETIGVQAIAMVTVPKRPVAPN

## SP059 nucleotide (SEQ ID NO:97)

CAACAGTCAGCTTCAGGAACGATTGAGGTGATTTACAGAGAAAAATGGCTCTGGGACACGGGGTGCCTT  
CACAGAAATCACAGGGATTCTCAAAAAAGACGGTGATAAAAAAATTGACAACACTGCCAAAACAGCTGT  
GATTCAAAATAGTACAGAAGGTGTTCTCTCAGCAGTTCAAGGGAATGCTAATGCTATCGGCTACATCTC  
CTTGGGATCTTTAACGAAATCTGTCAAGGCTTTAGAGATTGATGGTGTCAAGGCTAGTCGAGACACAGT  
TTTAGATGGTGAATACCCTCTTCAACGTCCCTTCAACATTGTTTGGTCTTCTAATCTTTCCAAGCTAGG  
TCAAGATTTTATCAGCTTTATCCACTCCAAACAAGGTCAACAAGTGGTCACAGATAATAAATTTATTGA  
AGCTAAAACCGAAACCACGGAATATACAAGCCAACACTTATCAGGCAAGTTGTCTGTTGTAGGTTCCAC  
TTCAGTATCTTCTTTAATGGAAAAATTAGCAGAAGCTTATAAAAAAGAAAATCCAGAAGTTACGATTGA  
TATTACCTCTAATGGGTCTTCAGCAGGTATTACCGCTGTTAAGGAGAAAACCGCTGATATTGGTATGGT  
TTCTAGGGAATTAACCTCTGAAGAAGGTAAGAGTCTCACCCATGATGCTATTGCTTTAGACGGTATTGC  
TGTTGTGGTCAATAATGACAATAAGGCAAGCCAAGTCAGTATGGCTGAACTTGACAGACGTTTTTTAGTGG  
CAAAATTAACCACCTGGGACAAGATTAA

Table 1

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**SP059 amino acid (SEQ ID NO:98)**

KQSASGTIEVISRENGSGTRGAFTEITGILKKDGDKKIDNTAKTAVIQNSTEGVLSAVQGNANAIGYIS  
LGS�TKSVKALEIDGVKASRDTVLDGEYPLQRPFNIVWSSNLSKLGQDFISFIHSKQGGQVVDNKFIE  
AKTETTEYTSQHLSGKLSVVGSTSVSSLMEKLAEAYKKENPEVTIDITSNGSSAGITAVKEKTADIGMV  
SRELTPEEGKSLTHDAIALDGIADVNNNDNKASQVSMALADVFSGKLTTWDKIK

**SP060 nucleotide (SEQ ID NO:99)**

ATTCGATGATGCGGATGAAAAGATGACCCGTGATGAAATTGCCTATATGCTGACAAATAGTGAAGAAAC  
ATTGGATGCTGATGAGATTGAGATGCTACAAGGTGTCTTTTCGCTCGATGAAGTGGCAGAGAGGT  
TATGGTTCCTCGAACGGATGCCTTTATGGTGGATATTCAGGATGATAGTCAAGCCATTATCCAAAGTAT  
TTTAAACAAAATTATTCCTCGTATCCCGGTTTATGATGGGGATAAGGACAATGTAATTGGAATCATTCA  
CACCAAGAGTCTCCTTAAGGCAGGCTTTGTGGACGGTTTGTGACAATATTGTTTGAAGAGAATTTTACA  
AGATCCACTTTTTGTACCTGAAACTATTTTTGTGGATGACTTGCTAAAAGAACTGCGAAATACCCAAAG  
ACAAATG

**SP060 amino acid (SEQ ID NO:100)**

FDDADEKMTREDEIAYMLTNSEETLDADEIEMLQGVFSLDELMAREVMVPRTD AFMVDIQDDSQAI IQSI  
LKQNSRIPVYDGDKNVIGIIHTKSLLKAGFVDGFDNIVWKRILQDPLFVPETIFVDDLLKELRNTQR  
QM

**SP062 nucleotide (SEQ ID NO:101)**

GGAGAGTCGATCAAAAGTAGATGAAGCTGTGTCTAAGTTTGAAAAGGACTCATCTTCTTCGTCAGTTTC  
AGACTCTTCCACTAAACCGGAAGCTTCAGATACAGCGAAGCCAAACAAGCCGACAGAACCAGGAGAAAA  
GGTAGCAGAAGCTAAGAAGAAGGTTGAAGAAGCTGAGAAAAAAGCCAAGGATCAAAAAGAAGAAGATCG  
TCGTAAC TACCAACCATTACTTACAAAACGCTTGAACTTGAAATTGCTGAGTCCGATGTGGAAGTTAA  
AAAAGCGGAGCTTGAAGTAGTAAAAGTGAAAGCTAACGAACCTCGAGACGAGCAA

**SP062 amino acid (SEQ ID NO:102)**

ESRSKVDEAVSKFEKDS SSSSSSSSSTKPEASDTAKPNKPTEPGEKVAEAKKKVEEA EKKAKDQKEEDR  
RNYPTITYKTLELEIAESDVEVKAELELVKVKANEPRDEQ

**SP063 nucleotide (SEQ ID NO:103)**

ATGGACAACAGGAAACTGGGACGAGGTTATATCTGGTAAGATTGACAAGTACAAAGATCCAGATATTCC  
AACAGTTGAATCACAAGAAGTTACGTCAGACTCTAGTGATAAAGAAATAACGGTAAGGTATGACCGTTT  
ATCAACACCAGAAAAACCAATCCCACAACCAATCCAGAGCATCCAAGTGTTCGACACCAAACCCAGA  
ACTACCAAATCAAGAGACTCCAACACCAGATAAACCAACTCCAGAACCAGGTACTCCAAAAACTGAAAC  
TCCAGTGAATCCAGACCCAGAAGTTCCGACTTATGAGACAGGTAAGAGAGAGGAATTGCCAAACACAGG  
TACAGAAGCTAAT

**SP063 amino acid (SEQ ID NO:104)**

WTTGNWDEVISGKIDKYKDPDIPTVESQEVTS DSSDKEITVRYDRLSTPEKPI PQPNPEHPSVPTPNPE  
LPNQETPTPKPTPEPGTPTKTETPVNPDPEVPTYETGKREELPNTGTEAN

**SP064 nucleotide (SEQ ID NO:105)**

CGATGGGCTCAATCCAACCCAGGTCAAGTCTTACCTGAAGAGACATCGGGAACGAAAGAGGGTGACTT  
ATCAGAAAAACCAGGAGACACCGTTCTCACTCAAGCGAAACCTGAGGGCGTTACTGGAAATACGAATTC  
ACTTCCGACACCTACAGAAAGAACTGAAGTGAGCGAGGAAACAAGCCCTTCTAGTCTGGATACACTTTT  
TGAAAAAGATGAAGAAGCTCAAAAAAATCCAGAGCTAACAGATGTCTTAAAAGAACTGTAGATACAGC  
TGATGTGGATGGGACACAAGCAAGTCCAGCAGAACTACTCCTGAACAAGTAAAAGGTGGAGTGAAAGA  
AAATACAAAAGACAGCATCGATGTTCTGCTGCTTATCTTGAAAAAGCTGAAGGGAAAGGTCTTTTAC  
TGCCGGTGTAACCAAGTAATTCCTTATGAACTATTCGCTGGTGATGGTATGTTAACTCGTCTATTACT  
AAAAGCTTCGGATAATGCTCCTTGGTCTGACAATGGTACTGCTAAAAATCCTGCTTTACCTCCTCTTGA  
AGGATTAACAAAAGGGAAATACTTCTATGAAGTAGACTTAAATGGCAATACTGTTGGTAAACAAGGTCA  
AGCTTTAATTGATCAACTTCGCGCTAATGGTACTCAAACCTTATAAAGCTACTGTAAAGTTTACGGAAA  
TAAAGACGGTAAAGCTGACTTGACTAATCTAGTTGCTACTAAAAATGTAGACATCAACATCAATGGATT  
AGTTGCTAAAGAAACAGTTCAAAAAGCCGTTGCAGACAACGTTAAAGACAGTATCGATGTTCCAGCAGC  
CTACCTAGAAAAAGCCAAGGGTGAAGGTCCATTACAGCAGGTGTCAACCATGTGATTCCATACGAACT  
CTTCGAGGTGATGGCATGTTGACTCGTCTCTTGCTCAAGGCATCTGACAAGGCACCATGGTCAGATAA

Table 1

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CGGCGACGCTAAAAACCCAGCCCTATCTCCACTAGGCGAAAACGTGAAGACCAAAGGTCAATACTTCTA  
TCAANTAGCCTTGGACGGAAATGTAGCTGGCAAAGAAAAACAAGCGCTCATTGACCAGTTCCGAGCAAA  
NGGTACTCAAACCTTACAGCGCTACAGTCAATGTCTATGGTAACAAAGACGGTAAACCAGACTTGGACAA  
CATCGTAGCAACTAAAAAAGTCACTATTAACATAAACGGTTTAATTTCTAAAGAAACAGTTCAAAAAGC  
CGTTGCAGACAACGTTAANGACAGTATCGATGTTCCAGCAGCCTACCTAGAAAAAGCCAAGGGTGAAGG  
TCCATTACAGCAGGTGTCAACCATGTGATTCCATACGAACCTCTTCGCAGGTGATGGTATGTTGACTCG  
TCTCTTGCTCAAGGCATCTGACAAGGCACCATGGTCAGATAACGGNGACGCTAAAAACCCAGCNCATC  
TCCACTAGGTGAAAACGTGAAGACCAAAGGTCAATACTTCTATCAANTAGCCTTGGACGGAAATGTAGC  
TGGCAAAGAAAAACAAGCGCTCATTGACCAGTTCCGAGCAAACGGTACTCAAACCTTACAGCGCTACAGT  
CAATGTCTATGGTAACAAAGACGGTAAACCAGACTTGGACAACATCGTAGCAACTAAAAAAGTCACTAT  
TAAGATAAATGTTAAAGAAACATCAGACACAGCAAATGGTTCATTATCACCTTCTAACTCTGGTTCCTGG  
CGTGACTIONGATGAATCACAATCATGCTACAGGTACTACAGATAGCATGCCTGCTGACACCATGACAAG  
TTCTACCAACACGATGGCAGGTGAAAACATGGCTGCTTCTGCTAACAAGATGTCTGATACGATGATGTC  
AGAGGATAAAGCTATG

**SP064 amino acid (SEQ ID NO:106)**

DGLNPTPGQVLPEETSGTKEGDLSEKPGDVTLTQAKPEGVTGNTNSLPTPTERTVEVSEETSPSSLDTLF  
EKDEEAQKNPELTDVLKETVDADVDGTQASPAETTPEQVKGKVKENTKDSIDVPAAYLEKAEGKGPFT  
AGVNQVPIPYELFAGDGMLTRLKLLKASDNAPWSDNGTAKNPALPPEGLTKGKYFYEVDLNGNTVKGQGO  
ALIDQLRANGTQTYKATVKVYGNKDGKADLTNLVATKNVDININGLVAKETVQKAVADNVKDSIDVPA  
YLEKAKGEGPFTAGVNHVPIPYELFAGDGMLTRLKLLKASDKAPWSDNGDAKNPALSPLGENVKTKGQYFY  
QXALDGNVAGKEKQALIDQFRAXGTQTYSATVNVYGNKDGKPDLDNIVATKKVTININGLISKETVQKA  
VADNVXDSIDVPAAYLEKAKGEGPFTAGVNHVPIPYELFAGDGMLTRLKLLKASDKAPWSDNGDAKNPALS  
PLGENVKTKGQYFYQXALDGNVAGKEKQALIDQFRANGTQTYSATVNVYGNKDGKPDLDNIVATKKVTI  
KINVKETSDTANGSLSPSNSGSGVTPMHNHATGTTDSMPADTMTSSNTMAGENMAASANKMSDTMMS  
EDKAM

**SP065 nucleotide (SEQ ID NO:107)**

TTCCAATCAAAAACAGGCAGATGGTAACTCAATATCGTGACAACCTTTTACCCTGTCTATGARTTTAC  
CAAGCAAGTCGCAGGAGATACGGCTAATGTAGAACTCCTAATCGGTGCTGGGACAGAACCTCATGAATA  
CGAACCATCTGCCAAGGCAGTTGCCAAAATCCAAGATGCAGATACCTTCGTTTATGAAAATGAAAACAT  
GGAAACATGGGTACCTAAATTGCTAGATACCTTGGATAAGAAAAAAGTGAAAACCATCAAGGCACAGG  
CGATATGTTGCTCTTGCCAGGTGGCGAGGAAGAAGAGGGAGACCATGACCATGGAGAAGAAGGTCATCA  
CCATGAGTTTGACCCCATGTTTGGTTATCACCAGTTTCGTGCCATtAACTAGTAGAGCACCATCCGCG  
ACACTTGTGACGAGATTATCCTGATAAAAAAGAGACCTTTGAGAAGAATGCAGCTGCCTATATCGAAAA  
ATTGCAAGCCTTGGATAAGGCTTACGCAGAAGGTTTGTCTCAAGCAAACAAAAGAGCTTTGTGACTCA  
ACACGCAgCCTTTAACTaTCTTGCCCTTGGACTATGGGACTC

**SP065 amino acid (SEQ ID NO:108)**

SNQKQADGKLNIVTTFYPVYEFTKQVAGDTANVELLIGAGTEPHEYEPSAKAVAKIQDADTFVYENENM  
ETWVPKLLDLDKKVKTIKATGDMLLLPGGEEEEGDHHDHGEEGHHHEFDPHVWLSPVRAIKLVEHHPR  
HLSADYPDKKETFEKNAAAYIEKLQALDKAYAEGLSQAKQKSFVTQHAAPNYLALDYGT

**SP067 nucleotide (SEQ ID NO:109)**

TATCACAGGATCGAACGGTAAGACAACCACAACGACTATGATTGGGGAAGTTTGTGACTGCTGCTGGCCA  
ACATGGTCTTTTATCAGGGAATATCGGCTATCCAGCTAGTCAGGTGCTCAAATAGCATCAGATAAGGA  
CAGCCTTGTTATGGAACCTTCTTCTTTCCAACCTCATGGGTGTTCAAGAATTCCATCCAGAGATTGCGGT  
TATTACCAACCTCATGCCAACTCATATCGACTACCATGGGTCAATTTTCGGAATATGTAGCAGCCAAGTG  
GAATATCCAGAACAAGATGACAGCAGCTGATTTCTTGTCTTGAACCTTTAATCAAGACTTGGCAAAAGA  
CTTGACTTCCAAGACAGAAGCCACTGTTGTACCATTTTCAACACTTGAAAAGGTTGATGGAGCTTATCT  
GGAAGATGGTCAACTCTACTTCCGTGGTGAAGTAGTCATGGCAGCGAATGAAATCGGTGTTCCAGGTAG  
CCACAATGTGGAATAATGCCCTTGGCAGCTATTGCTGTAGCCAAGCTTCGTGATGTGGACAATCAAACCAT  
CAAGGAAACTCTTTCAGCCTTCCGGTGGTGTCAAAACACCGTCTCCAGTTTGTGGATGACATCAAGGGTGT  
TAAATTCTATAACGACAGTAAATCAACTAATATCTTGGCTACTCAAAAAGCCTTGTGACGATTTGACAA  
CAGCAAGGTCGTCTTGATTGCAGGTGGTTTGGACCGTGGCAATGAGTTTGACGAATTGGTGCCAGACAT  
TACTGGACTCAAGAAGATGGTCATCTGGGTCAATCTGCAGAACGTGTCAAACGGGCAGCAGACAAGGC  
TGGTGTGCTTATGTGGAGGCGACAGATATTGCAGATGCGACCCGAAGGCCATGAGCTTGGGACTCA

Table 1

AGGAGATGTGGTTCTTCTTAGTCCTGCCAATGCTAGCTGGGATATGTATGCTAACTTTGAAGTACGTGG  
CGACCTCTTTATCGACACAGTAGCGGAGTTAAAAGAA

**SP067 amino acid (SEQ ID NO:110)**

GITGSNGKTTTTTMI GEVLTAAGQHLLSGNIGYPASQVAQIASDKDTLVMELSSFQLMGVQEFHPEIA  
VITNLMPTHIDYHGSFSEYVAAKWNIQNKMTAADFLVLNFNQDLAKDLTSKTEATVVPFSTLEKVDGAY  
LEDGQLYFRGEVMAANEIGVPGSHNVENALATIAVAKLRDNDQTIKETLSAFGGVKHRLQFVDDIKG  
VKFYNDKSKSTNILATQKALSGFDNSKVLIAGGLDRGNEFDELVPDITGLKKMVILGQSAERVKRAADK  
AGVAYVEATDIADATRKAYELATQGDVVLSPANASWDMYANFEVRGDLFIDTV AELKE

**SP068 nucleotide (SEQ ID NO:111)**

AAGTTCATCGAAGATGGTTGGGAAGTCCACTATATCGGGGACAAGTGTGGTATCGAACACCAAGAAATC  
CTTAAGTCAGGTTTGGATGTCACCTTCCATTCTATTGCGACTGGAAAATTGCGTCGCTATTTCTCTTGG  
CAAAATATGCTGGACGCTCTCAAAGTTGGTTGGGGAATTGTCCAATCGCTCTTTATCATGTTGCGACTG  
CGTCCACAGACCCTTTTTCAAAGGGGGGCTTTGTCTCAGTACCGCCTGTTATCGCTGCGCGTGTGTCA  
GGAGTGCCTGTCTTTATTACGAATCTGACCTGTCTATGGGCTTGGCCAATAAAATCGCCTATAAATTT  
CGGACTAAGATGTATTCAACCTTTGAACAAGCTTCGAGTTTGGCTAAGGTTGAGCATGTGGGAGCGG

**SP068 amino acid (SEQ ID NO:112)**

SSSKMVGKSTISGTSVVSNTKKSLSQVWMSPIILLRLENCVAISLGKICWTSSKLVGELSNRSLSCDC  
VHRPFFQRGALSQYRLLSLRVCQECLSLFTNLTLCLWAWPIKSPINLRLRCIQPLNKL RVWLRLSMWER

**SP069 nucleotide (SEQ ID NO:113)**

ATCGCTAGCTAGTGAAATGCAAGAAAGTACACGTAAATTCAAGGTTACTGCTGACCTAACAGATGCCGG  
TGTTGGAACGATTGAAGTTCCTTTGAGCATTGAAGATTTACCCAATGGGCTGACCGCTGTGGCGACTCC  
GCAAAAATTACAGTCAAGATTGGTAAGAAGGCTCAGAAGGATAAGGTAAAGATTGTACCAGAGATTGA  
CCCTAGTCAAATTGATAGTCGGGTACAAATTGAAATGTCTATGGTGTCTAGATAAAGAAGTGTCTATTAC  
GAGTGACCAAGAGACATTGGATAGAATTGATAAGATTATCGCTGTTTTGCCAACTAGCGAACGTATAAC  
AGGTAATTACAGTGGTTTCAGTACCTTTGCAGGCAATCGACCGCAATGGTGTGTCTTACCGGCAGTTAT  
CACTCCGTTTGATACAATAATGAAGGTGACTACAAAACCAGTAGCACCAAGTTCAAGCACATCAAATTC  
AAGTACAAGCAGTTCATCGGAGACATCTTCGTCAACGAAAGCAACTAGTTCAAAAACGAAT

**SP069 amino acid (SEQ ID NO:114)**

SLASEMQESTRKFKVTADLTDAGVGTIEVPLSIEDLPNGLTAVATPQKITVKIGKKAQKDKVKIVPEID  
PSQIDSRVQIENVMVSDKEVSITSQDETLDRIKIIAVLPTSERITGNYSVPLQAI DRNGVVLPAVI  
TPFDTIMKVTTKPVAPSSSTSNSSTSSSSETSSSTKATSSKTN

**SP070 nucleotide (SEQ ID NO:115)**

GCACCAGATGGGGCACAAGGTTTCAGGGATCAGATGTTGAAAAGTACTACTTTACCCAACGCGGTCTTGA  
GCAGGCAGGAATTACCATTCTTCTTTTGATGAAAAAATCTAGACGGTGATATGGAAATTATCGCTGG  
AAATGCCTTTCGTCCAGATAACAACGTCGAAATTGCCTATGCGGACCAAAATGGTATCAGCTACAAACG  
TTACCATGAGTTTCTAGGTAGCTTTATGCGTGACTTTGTTAGCATGGGAGTAGCAGGAGCACATGGAAA  
AACTTCAACGACAGGTATGTTGTCTCATGTCTTGTCTCACATTACAGATACCAGCTTCTTGATTGGAGA  
TGGGACAGGTCGTGGTTTCGGCCAATGCCAAATATTTTGTCTTTGAATCTGACGAATATGAGCGTCACTT  
CATGCCCTTACCACCCAGAATACTCTATTATCACCAACATTGACTTTGACCATCCAGATTATTTACAAG  
TCTCGAGGATGTTTTTAATGCCTTTAACGACTATGCCAAACAAATCACCAGGGTCTTTTGTCTATGG  
TGAAGATGCTGAATTGCGTAAGATTACGTCGTATGCACCAATTTATTATTATGGTTTTGAAGCTGAAGG  
CAATGACTTTGTAGCTAGTGATCTTCTTCGTTCAATAACTGGTTCAACCTTCACCGTTTCAATTTCCGTGG  
ACAAAATTTGGGGCAATTCCACATTCCAACCTTTGGTCGTCACAATATCATGAATGCGACAGCCGTTAT  
TGGTCTTCTTTACACAGCAGGATTTGATTTGAACTTGGTGCGTGAGCACTTGAAAACATTTGCCGGTGT  
TAAACGTCGTTTCACTGAGAAAATTGTCAATGATACAGTGATTATCGATGACTTTGCCACCATCCAAC  
AGAAATTATTGCGACCTTGGATGCGGCTCGTCAGAAATACCCAAGCAAGGAAATTGTAGCAGTCTTTCA  
ACCGCATACCTTTACAAGAACCATTGCCTTGTGGACGACTTTGCCCATGCTTTAAACCAAGCAGATGC  
TGTTTATCTAGCGCAAATTTATGGCTCGGCTCGTGAAGTAGATCATGGTGACGTTAAGGTAGAAGACCT  
AGCCAACAAAATCAACAAAAACACCAAGTGATTACTGTTGAAAATGTTTCTCCACTCCTAGACCATGA  
CAATGCTGTTTACGTCTTTATGGGAGCAGGAGACATCCAAACCTATGAATACTCATTTGAGCGTCTCTT  
GTCTAACTTGACAAGCAATGTTCAA

Table 1

## SP070 amino acid (SEQ ID NO:116)

HQMGHKVQGS DVEKYFTQ RGLEQAGITILPFDEKNLDGDM EIIAGNAFRPD MNVEIAYADQNGISYKR  
YHEFLGSFMRDFVSMGVAGAHGKTSTTGMLSHVLSHITDTSFLIGDGTGRGSANAKYFVFESDEYERHF  
MPYHPEYSIITNIDFDHPDYFTSLEDVFNAFNDYAKQITKGLFVYGEDAELRKITS DAPIYYYGF EAEG  
NDFVASDLLRSITGSTFTVHFRGQNLGQFHIPTFGRHNIMNATAVIGLLYTAGFDLNLVREHLKTFAGV  
KRRETEKIVNDTVIIDDFAHHPTEIIATLDAARQKYP SKEIVAVFQPHFTFTRTIALLDFAHALNQADA  
VYLAQIYGSAREVDHGDVKVEDLAN KINKKHQVITVENVSPLLDHDNAVYVFMGAGDIQTYEYSFERLL  
SNLTSNVQ

## SP071 nucleotide (SEQ ID NO:117)

TTTAAACCCAACTGTTGGTACTTTTCCTTTTACTGCAGGATTGAGCTTGTTAGT TTTATTGGTTTCTAA  
AAGGGAAATGGAAGAAACGACTTGTTCATTTTCTGCTGTTGACTAGCATGGGAGTTCAATTGTTGCC  
GGCCAGTGCTTTTGGGTTGACCAGCCAGATTTTATCTGCCTATAATAGTCAGCTTTCTATCGGAGTCGG  
GGAACATTTACCAGAGCCTCTGAAAATCGAAGGTTATCAATATATTGGTTATATCAAAACTAAGAAACA  
GGATAATACAGAGCTTTCAAGGACAGTTGATGGGAAATACTCTGCTCAAAGAGATAGTCAACCAAACCTC  
TACAAAACATCAGATGTAGTTCATT CAGCTGATTTAGAATGGAACCAAGGACAGGGGAAGGTTAGTTT  
ACAAGGTGAAGCATCAGGGGATGATGGACTTT CAGAAAAATCTTCTATAGCAGCAGACAATCTATCTTC  
TAATGATT CATTTCGCAAGTCAAGTTGAGCAGAATCCGGATCACAAGGAGAATCTGTAGTTCGACCAAC  
AGTGCCAGAACAAGGAAATCCTGTGTCTGCTACAACGGTGCAGAGTGCGGAAGAGGAAGTATTGGCGAC  
GACAAATGATCGACCAGAGTATAAACTTCCATTGGAAACCAAAGGCACGCAAGAACCCGGTCATGAGGG  
TGAAGCCGCAGTCCGTGAAGACTTACCAGTCTACACTAAGCCACTAGAAACCAAAGGTACACAAGGACC  
CGGACATGAAGGTGAAGCTGCAGTTCCGCGAGGAAGAACCAGCTTACACAGAACC GTTAGCAACGAAAGG  
CACGCAAGAGCCAGGTCATGAGGGCAAAGCTACAGTCCGCGAAGAGACTCTAGAGTACACGGAACCGGT  
AGCGACAAAAGGCACACAAGAACC CGAACATGAGGGCGAaCGGsCAGTAGAAGAAGAATTC CGGCTTT  
AGAGGTCACTACACGAAATAGAACGAAATCCAGAATATTCCTTATACAACAGAAGAAATTCAGGATCC  
AACACTTCTGAAAAATCGTCGTAAGATTGAACGACAAGGGCAAGCAGGGACACGTACAATTCAATATGA  
AGACTACATCGTAAATGGTAAATGTCGTAGAACTAAAGAAGTGT CACGAATGAAGTAGCTCCGGTCAA  
CGAAGTCGTTAAAGTAGGAACACTTGTGAAAGTTAAAGCTACAGTAGAAATTACAACTTAACAAAAGT  
TGAGAACAAAAATCTATAACTGTAAGTTATAACTTAATAGACACTACCTCAGCATATGTTTCTGCAAA  
AACGCAAGTTTTCCATGGAGACAAGCTAGTTAAAGAGGTGGATATAGAAAATCCTGCCAAAGAGCAAGT  
AATATCAGGTTT TAGATTACTACACACCGTATACAGTTAAAACACACCTAAC TTATAATTTGGGTGAAAA  
TAATGAGGAAAATACTGAAACATCAACTCAAGATTTCCAATTAGAGTATAAGAAAATAGAGATTAAAGA  
TATTGATT CAGTAGAATTATACGGTAAAGAAAATGATCGTTATCGTAGATATTTAAGTCTAAGTGAAGC  
GCCGACTGATACGGCTAAATACTTTGTAAAAGTGAAATCAGATCGCTTCAAAGAAATGTACCTACCTGT  
AAAATCTATTACAGAAAATACGGATGGAACGTATAAAGTGACGGTAGCCGTTGATCAACTTGTGCAAGA  
AGGTACAGACGGTTACAAAGATGATTACACATTTACTGTAGCTAAATCTAAAGCAGAGCAACCAGGAGT  
TTACACATCCTTTAAACAGCTGGTAACAGCCATGCAAAGCAATCTGTCTGGTGTCTATACATTGGCTTC  
AGATATGACCGCAGATGAGGTGAGCTTAGGCGATAAGCAGACAAGTTATCTCACAGGTGCATTTACAGG  
GAGCTTGATCGGTTCTGATGGAACAAAATCGTATGCCATTTATGATTTGAAGAAACCATTATTTGATAC  
ATTAAATGGTGCTACAGTTAGAGATTTGGATATTA AAAC TGTTC TGCTGATAGTAAAGAAAATGTGCG  
AGCGCTGGCGAAGGCAGCGAATAGCGCGAATATTAATAATGTTGCAGTAGAAGGAAAAATCTCAGGTGC  
GAAATCTGTTGCGGGATTAGTAGCGAGCGCAACAAATACAGTGATAGAAAACAGCTCGTTTACAGGGAA  
ACTTATCGCAAATACCCAGGACAGTAATAAAAATGATAC TGGAGGAATAGTAGGTAATATAACAGGAAA  
TAGTTTCGAGAGTTAATAAAGTTAGGGTAGATGCCTTAATCTCTACTAATGCACGCAATAATAACCAAAC  
AGCTGGAGGGATAGTAGGTAGATTAGAAAATGGTGCAATTGATATCTAATTCGTTGCTACTGGAGAAAT  
ACGAAATGGTCAAGGATATTCTAGAGTCGGAGGAATAGTAGGATCTACGTGGCAAAACGGTCGAGTAA  
TAATGTTGTGAGTAACGTAGATGTTGGAGATGGTTATGTTATCACC GG TGATCAATACGCAGCAGCAGA  
TGTGAAAAATGCAAGTACATCAGTTGATAATAGAAAAGCAGACAGATTTCGCTACAAAATTATCAAAAGA  
CCAAATAGACGCGAAAGTTGCTGATTATGGAATCACAGTAACTCTTGATGATACTGGGCAAGATTTAAA  
ACGTAATCTAAGAGAAGTTGATTATACAAGACTAAATAAAGCAGAAGCTGAAAGAAAAGTAGCTTATAG  
CAACATAGAAAAACTGATGCCATTCTACAATAAAGACCTAGTAGTTCACTATGGTAACAAAGTAGCGAC  
AACAGATAAACTTTACTACTACAGAATTGTTAGATGTTGTGCCGATGAAAGATGATGAAGTAGTAACGGA  
TATTAATAATAAGAAAAATTCAATAAATAAAGTTATGTTACATTTCAAAGATAATACAGTAGAATACCT  
AGATGTAACATTC AAAGAAAAC TTCATAAACAGTCAAGTAATCGAATACAATGTTACAGGAAAAGAATA  
TATATTCACACCAGAAGCATTGTTTTCAGACTATACAGCGATAACGAATAACGTACTAAGCGACTTGCA  
AAATGTAACACTTAAC

## SP071 amino acid (SEQ ID NO:118)

Table 1

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FNPTVGTFLFTAGLSLLVLLVSKRENGKKRLVHFLLLTSMGVQLLPASAFGLTSQILSAYNSQLSIGVG  
 EHLPEPLKIEGYQYIGYIKTKKQDNTELSRTVDGKYSAQRDSQPNSTKTSADVHSADLEWNQGGKVS  
 QGEASGDDGLSEKSSIAADNLSSNDSFASQVEQNPDPKHGESVVRPTVPEQGNPVSATTVQSAEEEEVLAT  
 TNDREPKLPLETKGTQEPGHEGEAAVREDLPVYTKPLETKGTQGPGEHEGEAAVREEEPAYTEPLATKG  
 TQEPGHEGKATVREETLEYTEPVATKGTQEPHEGERXVEEELPALEVTTRNRTEIQNIPYTTEEIQDP  
 TLLKNRRKIERQGGAGTRTIQYEDYIVNGNVVETKEVSRTEVAPVNEVVKVGLTVKVKPTVEITNLTKV  
 ENKKSITVSYNLIDTTSAYVSAKTQVFHGDKLVEVDIENPAKEQVISGLDYYTPYTVKTHLTYNLGEN  
 NEENTETSTQDFQLEYKKIEIKDIDSVELYKENDRYRRYLSLSEAPTD TAKYFVKVKSDFKEMYL PV  
 KSITENTDGTYKVTVAVDQLVEEGTDGYKDDYTFTVAKSKAEQPGVYTSFKQLVTAMQSNLSGVYTLAS  
 DMTADEVSLGDKQTSYLTGAFTGSLIGSDGTSYAIYDLKKPLFDTLNGATVRDLDIKTVSADSKENVA  
 ALAKAANSANINNVAVEGKISGAKSVAGLVSATNTVIENSSFTGKLIANHQDSNKNNDTGIVGNITGN  
 SSRVKNVRVDALISTNARNNNQTAGGIVGRLENGALISNSVATGEIRNGQGYSRVGGIVGSTWQNGRVN  
 NVVSNVDVGDGYVITGDQYAAADVKNASTSVDNRKADRFATKLSKDQIDAKVADYGITVTLDDTGQDLK  
 RNLREVDTYRLNKAERKVAYSNIEKLMPFYNKDLVVHYGNKVATTDKLYTTELLDVVPMKDDEVVTD  
 INNKKNSINKVMLHFKDNTVEYLDVTFKENFINSQVIEYNVTGKEYIFTPEAFVSDYTAITNNVLSDLQ  
 NVTNLN

**SP072 nucleotide (SEQ ID NO:119)**

TTTTAACCCAACGTGTGGTACTTTCTCTTTTACTGCAGGATTGAGCTTGTTAGTTTTATTGGTTTCTAA  
 AAGGGAAAATGGAAGAAACGACTTGTTCATTTTCTGCTGTTGACTAGCATGGGAGTTCAATTGTTGCC  
 GGCCAGTGCTTTTGGGTGACCAGCCAGATTTTATCTGCCTATAATAGTCAGCTTTCCTATCGGAGTCGG  
 GGAACATTTACCAGAGCCTCTGAAAATCGAAGGTTATCAATATATTGGTTATATCAAACTAAGAAAACA  
 GGATAATACAGAGCTTTCAGGACAGTTGATGGGAAATACTCTGCTCAAAGAGATAGTCAACCAAACCTC  
 TACAAAACATCAGATGTAGTTTACCTCAGCTGATTTAGAATGGAACCAAGGACAGGGGAAGGTTAGTTT  
 ACAAGGTGAAGCATCAGGGGATGATGGACTTTCAGAAAATCTTCTATAGCAGCAGACAATCTATCTTC  
 TAATGATTCATTCGCAAGTCAAGTTGAGCAGAATCCGGATCACAAAGGAGAATCTGTAGTTCGACCAAC  
 AGTGCCAGAACAAGGAAATCCTGTGTCTGCTACAACGGTGCAGAGTGCAGGAGAGGAAGTATTGGCGAC  
 GACAAATGATCGACCAGAGTATAAACTTCCATTGGAACCAAAGGCACGCAAGAACCCGGTCATGAGGG  
 TGAAGCCGAGTCCGTGAAGACTTACAGTCTACACTAAGCCACTAGAAACCAAAGGTACACAAGGACC  
 CGGACATGAAGGTGAAGCTGCAGTTCGCGAGGAAGAACCAGCTTACACAGAACCGTTAGCAACGAAAGG  
 CACGCAAGAGCCAGGTCATGAGGGCAAAGCTACAGTCCGCGAAGAGACTCTAGAGTACACGGAACCGGT  
 AGCGACAAAAGGCACACAAGAACCCGAACATGAGGGCGAaCGGsCAGTAGAAGAAGAACTTCCGGCTTT  
 AGAGGTCACTACACGAAATAGAACGGAATCCAGAATATTCCTTATACAACAGAAGAAATTCAGGATCC  
 AACACTTCTGAAAAATCGTCGTAAGATTGAACGACAAGGGCAAGCAGGGACACGTACAATTCAATATGA  
 AGACTACATCGTAAATGGTAATGTCGTAGAACTAAAGAAGTGTACGAACTGAAGTAGCTCCGGTCAA  
 CGAAGTCGTTAAAGTAGGAACACTTGTGAAAGTTAAACCTACAGTAGAAATTACAACTTAACAAAAGT  
 TGAGAACAAAAAATCTATAACTGTAAGTTATACTTAATAGACACTACCTCAGCATATGTTTCTGCAAA  
 AACGCAAGTTTCCATGGAGACAAGCTAGTTAAAGAGGTGGATATAGAAAAATCCTGCCAAAGAGCAAGT  
 AATATCAGGTTTAGATTACTACACACCGTATACAGTTAAACACACCTAACTTATAATTTGGGTGAAAA  
 TAATGAGGAAAATACTGAAACATCAACTCAAGATTTCCAATTAGAGTATAAGAAAATAGAGATTAAAGA  
 TATTGATTCAGTAGAATTATACGGTAAAGAAAATGATCGTTATCGTAGA

**SP072 amino acid (SEQ ID NO:120)**

FNPTVGTFLFTAGLSLLVLLVSKRENGKKRLVHFLLLTSMGVQLLPASAFGLTSQILSAYNSQLSIGVG  
 EHLPEPLKIEGYQYIGYIKTKKQDNTELSRTVDGKYSAQRDSQPNSTKTSADVHSADLEWNQGGKVS  
 QGEASGDDGLSEKSSIAADNLSSNDSFASQVEQNPDPKHGESVVRPTVPEQGNPVSATTVQSAEEEEVLAT  
 TNDREPKLPLETKGTQEPGHEGEAAVREDLPVYTKPLETKGTQGPGEHEGEAAVREEEPAYTEPLATKG  
 TQEPGHEGKATVREETLEYTEPVATKGTQEPHEGERXVEEELPALEVTTRNRTEIQNIPYTTEEIQDP  
 TLLKNRRKIERQGGAGTRTIQYEDYIVNGNVVETKEVSRTEVAPVNEVVKVGLTVKVKPTVEITNLTKV  
 ENKKSITVSYNLIDTTSAYVSAKTQVFHGDKLVEVDIENPAKEQVISGLDYYTPYTVKTHLTYNLGEN  
 NEENTETSTQDFQLEYKKIEIKDIDSVELYKENDRYRR

**SP073 nucleotide (SEQ ID NO:121)**

TCGTAGATATTTAAGTCTAAGTGAAGCGCCGACTGATACGGCTAAATACTTTGTAAAAGTGAAATCAGA  
 TCGCTTCAAAGAAATGTACCTACCTGTAAATCTATTACAGAAAAATACGGATGGAACGTATAAAGTGAC  
 GGTAGCCGTTGATCAACTTGTGCAAGAAGGTACAGACGGTTACAAAGATGATTACACATTTACTGTAGC  
 TAAATCTAAAGCAGAGCAACCAGGAGTTTACACATCCTTTAAACAGCTGGTAACAGCCATGCAAGCAA  
 TCTGTCTGGTGTCTATACATTGGCTTCAGATATGACCGCAGATGAGGTGAGCTTAGGCGATAAGCAGAC

Table 1

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AAGTTATCTCACAGGTGCATTTACAGGGAGCTTGATCGGTTCTGATGGAACAAAATCGTATGCCATTTA  
TGATTGAAGAAACCATTATTTGATACATTAAATGGTGCTACAGTTAGAGATTGGATATTAACACTGT  
TTCTGCTGATAGTAAAGAAAATGTCGCAGCGCTGGCGAAGGCAGCGAATAGCGCGAATATTAATAATGT  
TGCAGTAGAAGGAAAAATCTCAGGTGCGAAAATCTGTTGCGGGATTAGTAGCGAGCGCAACAAATACAGT  
GATAGAAAAACAGCTCGTTTACAGGGAACCTTATCGCAAATCACCAGGACAGTAATAAAAATGATACTGG  
AGGAATAGTAGGTAATATAACAGGAAATAGTTCGAGAGTTAATAAAGTTAGGGTAGATGCCTTAATCTC  
TACTAATGCACGCAATAATAACCAAACAGCTGGAGGGATAGTAGGTAGATTAGAAAAATGGTGCATTGAT  
ATCTAATTCGGTTGCTACTGGAGAAATACGAAATGGTCAAGGATATTCTAGAGTCGGAGGAATAGTAGG  
ATCTACGTGGCAAAACGGTCGAGTAAATAATGTTGTGAGTAACGTAGATGTTGGAGATGGTTATGTTAT  
CACCGGTGATCAATACGCAGCAGCAGATGTGAAAAATGCAAGTACATCAGTTGATAATAGAAAAGCAGA  
CAGATTCGCTACAAAATTATCAAAAGACCAAATAGACGCGAAAGTTGCTGATTATGGAATCACAGTAAC  
TCTTGATGATACTGGGCAAGATTTAAACGTAATCTAAGAGAAGTTGATTATACAAGACTAAATAAAGC  
AGAAGCTGAAAGAAAAGTAGCTTATAGCAACATAGAAAAACTGATGCCATTCTACAATAAAGACCTAGT  
AGTTCACCTATGGTAAACAAAGTAGCGACAACAGATAAACTTTACACTACAGAATTGTTAGATTGTTGTGCC  
GATGAAAGATGATGAAGTAGTAACGGATATTAATAATAAGAAAAATTCAATAAATAAAGTTATGTTACA  
TTTCAAAGATAATACAGTAGAATACCTAGATGTAACATTCAAAGAAAACCTCATAAACAGTCAAGTAAT  
CGAATACAATGTTACAGGAAAAGAATATATATTACACCAGAAGCATTGTTTCAGACTATACAGCGAT  
AACGAATAACGTACTAAGCGACTTGCAAAATGTAACACTTAAC

**SP073 amino acid (SEQ ID NO:122)**

RRYLSLSEAPDITAKYFVKVKSDFKEMYLVPKSI TENDGTYKVTVAVDQLVEEGTDGYKDDYFTVA  
KSKAEQPGVYTSFKQLVTAMQSNLSGVYTLASDMTAEVSLGDKQTSYLTGAFTGSLIGSDGTSYAIY  
DLKKPLFDTLNGATVRDLDIKTVSADSKENVAALAKAANSANINNVAVEGKISGAKSVAGLVASATNTV  
IENSSFTGKLIANHQDSNKNDTGGIVGNITGNSSRVNKVRDALISTNARNNNQTAGGIVGRLENGALI  
SNSVATGEIRNGQGYSRVGGIVGSTWQNGRVNNVSNVDVGDGYVITGDQYAAADVKNASTSVDNRKAD  
RFATKLSKDQIDAKVADYGITVTLDDTGQDLKRNLEVDYTRLNKAERKVAYSNIEKLMPFYNKDLV  
VHYGNKVATTDKLYTTELLDVVPMKDDEVVTDINNKNKSNKVMLEHFKDNTVEYLDVTFKENFINSQVI  
EYNVTGKEYIFTPEAFVSDYTAITNNVLSLDLQNVTLN

**SP074 nucleotide (SEQ ID NO:123)**

CTTTGGTTTTGAAGGAAGTAAGCGTGGACAATTTGCTGTAGAAGGAATCAATCAACTTCGTGAGCATGT  
AGACACTCTATTGATTATCTCAAACAACAATTTGCTTGAAATTGTTGATAAGAAAACACCGCTTTTGGA  
GGCTCTTAGCGAAGCGGATAACGTTCTTCGTCAAGGTGTTCAAGGGATTACCGATTTGATTACCAATCC  
AGGATTGATTAACCTTGACTTTGCCGATGTGAAAACGGTAATGGCAAACAAAGGGAATGCTCTTATGGG  
TATTGGTATCGGTAGTGGAGAAGAACGTGTGGTAGAAGCGGCACGTAAGGCAATCTATTCACCACTTCT  
TGAAACAACCTATTGACGGTGTGAGGATGTTATCGTCAACGTTACTGGTGGTCTTGACTTAACCTTGAT  
TGAGGCAGAAGAGGCTTCACAAATTGTGAACCAGGCAGCAGGTCAAGGAGTGAACATCTGGCTCGGTAC  
TTCAATTGATGAAAGTATGCGTGATGAAATTCGTGTAACAGTTGTTGCAACGGGTGTTTCGTCAAGACCG  
CGTAGAAAAGGTTGTGGCTCCACAAGCTAGATCTGCTACTAATACTACCGTGAGACAGTGAAACCAGCTCA  
TTCACATGGCTTTGATCGTCATTTTGATATGGCAGAAACAGTTGAATTGCCAAAACAAAATCCACGTCTG  
TTTGAACCAACTCAGGCATCTGCTTTTGGTGATTGGGATCTTCGCCGTGAATCGATTGTTTCGTACAAC  
AGATTCAGTCGTTTCTCCAGTCGAGCGCTTTGAAGCCCCAATTTACAAGATGAAGATGAATTGGATAC  
ACCTCCATTTTTCAAAAATCGT

**SP074 amino acid (SEQ ID NO:124)**

FGFEGSKRGQFAVEGINQLREHVDTLIIISNNLLEIVDKKTPLEALSEADNVLRQGVQGITDLITNP  
GLINLDFADVKTVMANKGNALMGIGIGSGEERVVEAARKAIYSPLETTIDGAEDVIVNVTTGGLDLTLI  
EAEASQIVNQAGQGVNIWLGTSIDESMRDEIRVTVVATGVRQDRVEKVVPQARSATNYRETVKPAH  
SHGFDRHFDMAETVELPKQNPRLPTQSAFGDWDLRRESIVRTTDSVVSVERFEAPISQDEDELDT  
PPFFKNR

**SP075 nucleotide (SEQ ID NO:125)**

CTACTACCTCTCGAGAGAAAGTGACCTAGAGGTGACCGTTTTTGACCATGAGCAAGGTCAAGCCACCAA  
GGCCGCAGCAGGAATTATCAGTCCTTGTTTTCCAAACGCCGTAATAAAGCCTGGTACAAGATGGCGCG  
CTTGGGGGCTGATTTTTATGTGGATTTATTAGCTGATTTAGAGAAATCAGGACAAGAAATCGACTTTTA  
CCAGCGTTCCGGAGTCTTTCTCTTGAAAAGGATGAATCCAATTTGGAAGAACTTTATCAACTGGCCCT  
CCAGCGCAGAGAAGAATCTCCCTTGATAGGGCAATTAGCCATTCTGAACCAAGCCTCAGCTAATGAATT  
ATTCCTGTTTGCAGGGATTTGACCGCTGCTCTATGCTTCTGGTGGAGCGAGAGTAGATGGCCAAC

Table 1

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TTTAGTGACTCGTTTGTCTGGAAGTCAGTTCATGTCAAGCTGGTCAAAGAAAAAGTGACTCTGACACCGTT  
AGCATCAGGCTACCAGATTGGTGAAGAGGAGTTTGTAGCAGGTTATTTTGGCGACGGGAGCTTGGTTGGG  
GGACATGTTAGAGCCTTTAGGTTATGAAGTGGATGTCCGTCCTCAAAAAGGACAACCTACGAGATTATCA  
GCTTGCCCAAGACATGGAAGATTACCCTGTTGTCATGCCAGAAGGGGAGTGGGATTTGATTCCCTTTGC  
AGGTGGGAAATTATCCTTAGGCGCTACCCACGAAAATGACATGGGATTTGATTTGACGGTAGATGAAAC  
CTTGCTCCAACAAATGGAGGAGGCCACCTTGACTCACTATCTGATTTTGGCTGAAGCTACTTCAAAATC  
TGAGCGTGTGGAATCCGTGCCCTACACCAGTGATTTCTCTCCTTTCTTTGGGCAGGTGCCTGACTTAAC  
TGGTGTCTATGCAGCCAGTGGACTAGGTTTCATCAGGCCTCACAACTGGTCTTATCATTGGTTACCATCT  
AGCCCAACTGATCCAAGACAAGGAGTTGACCTTGGACCTCTAAATTACCCAATTGAAAACCTATGTCAA  
ACGAGTAAAAAGCGAA

**SP075 amino acid (SEQ ID NO:126)**

YYLSRESLEVTVDHEQGGATKAAAGIISPWFSSKRRNKAWYKMARLGADFYVDLLADLEKSGQEIDFY  
QRSGVFLKKDESNLEELYQLALQRREESPLIGQLAILNQASANELFPGLQGFDRLLYASGGARVDGQL  
LVTRLLEVSHVKLVKEKVTLTPLASGYQIGEEFEQVILATGAWLGDMLEPLGYEVDVRPQKGQLRDYQ  
LAQDMEDYPVVMPEGEWDLIPFAGGKLSLGATHENDMGFDLTVDLTLQOMEEATLTHYLILAEATSKS  
ERVGIRAYTSDFPSFFGQVDPDLTGVAASGLGSSGLTTGPIIGYHLAQLIQDKELTLDPLNPIENYVK  
RVKSE

**SP076 nucleotide (SEQ ID NO:127)**

TAAGGTCAAAAGTCAGACCGCTAAGAAAGTGCTAGAAAAGATTGGAGCTGACTCGGTTATCTCGCCAGA  
GTATGAAATGGGGCAGTCTCTAGCACAGACCATTCTTTTCCATAATAGTGTTGATGTCTTTTCAGTTGGA  
TAAAAATGTGTCTATCGTGGAGATGAAAATTCCTCAGTCTTGGGCAGGTCAAAGTCTGAGTAAATTAGA  
CCTCCGTGGCAAATACAATCTGAATATTTTGGGTTTCCGAGAGCAGGAAAATTCCTTGGATGTTGA  
ATTTGGACCAGATGACCTCTTGAAAGCAGATACCTATATTTTGGCAGTCATCAACAACAGTATTTGGA  
TACCTA

**SP076 amino acid (SEQ ID NO:128)**

KVKSQTAKKVLEKIGADSVISPEYEMGQSLAQITLFHNSVDVFQLDKNVSIIVEMKIPQSWAGQSLSKLD  
LRGKYNLNLGFRQENSPLDVEFGPDDLKADTYILAVINNQYLDTL

**SP077 nucleotide (SEQ ID NO:129)**

TGACGGGTCTCAGGATCAGACTCAGGAAATCGCTGAGTGTCTTAGCTAGCAAGTATCCTAATATCGTTAG  
AGCCATCTATCAGGAAAATAAATGCCATGGCGGTGCGGTCAATCGTGGCTTGGTAGAGGCTTCTGGGCG  
CTATTTTAAAGTAGTTGACAGTGATGACTGGGTGGATCCTCGTGCCTACTTGAAAATCTTGAAACTTG  
CAGGAACTTGAGAGCAAAGGTCAAGAGGTGGATGTCTTTG

**SP077 amino acid (SEQ ID NO:130)**

DGSQDQTQEIACLASKYPNIVRAIYQENKCHGGA VNRGLVEASGRYFKVVDSDDWVDPAYL KILETC  
RNLRAKVKRWMSL

**SP078 nucleotide (SEQ ID NO:131)**

TAGAGGCTTTGCCAAATGGTGGGAAGGGCACGAGCGTCGAAAAGAGGAACGCTTTGTCAAACAAGAAGA  
AAAAGCTCGCCAAAAGGCTGAGAAAGAGGCTAGATTAGAACAAGAAGAGACTGAAAAAGCCTTACTCGA  
TTTGCCTCCTGTTGATATGGAACGGGTGAAATCTGACAGAGGAAGCTGTTCAAATCTTCCACCTAT  
TCCAGAAGAAAAGTGGGTGGAACCAGAAATCATCCTGCCTCAAGCTGAACCTAAATTCCTGAACAGGA  
AGATGACTCAGATGACGAAGATGTTTCAGGTCGATTTTTCAGCCAAAGAAGCCCTTGAATACAACTTCC  
AAGCTTACAACCTTTTGCACCAGATAAACCAAAAGATCAGTCTAAAGAGAAGAAAATTGTCAGAGAAAA  
TATCAAATCTTTAGAAGCAACCTTTGCTAGCTTTGGTATTAAGGTAACAGTTGAACGGGCGGAAATTGG  
GCCATCAGTGACCAAGTATGAAGTCAAGCCGGCTGTTGGTGTAAGGGTCAACCGCATTTCCAATCTATC  
AGATGACCTCGCTCTAGCCTTGGCTGCCAAAGATGTCCGGATTGAAGCACCAATCCCTGGGAAATCCCT  
AATCGGAATTGAAGTGCCCAACTCCGATATTGCCACTGTATCTTTCCGAGAACTATGGGAACAATCGCA  
AACGAAAGCAGAAAATTTCTTGGAAATTCCTTTAGGGAAGGCTGTTAATGGAACCGCAAGAGCTTTTGA  
CCTTTCTAAAATGCCCCACTTGCTAGTTGCAGGTTCAACGGGTTCAAGGAAGTCAGTAGCAGTTAACGG  
CATTATTGCTAGCATTCTCATGAAGGCGAGACCAGATCAAGTTAAATTTATGATGGTTCGATCCCAAGAT  
GGTTGAGTTATCTGTTTACAATGATATCCCCACCTCTTGATTCCAGTCGTGACCAATCCACGCAAAGC  
CAGCAAGGCTCTGCAAAGGTTGTGGATGAAATGGAAAACCGTTATGAACTCTTTGCCAAGGTGGGAGT  
TCGGAATATTGCAGGTTTTAATGCCAAGGTAGAAGAGTTCAATCCCAGTCTGAGTACAAGCAAATTC



Table 1

GCTACCATTTCATTGTCGTGATTGTGGATGAGTTGGCTGACCTCATGATGGTGGCCAGCAAGGAAGTGGA  
AGATGCTATCATCCGCTCTTGGGCAGAAAGGCGCGTGTGCAAGGTATCCACATGATTCTTGCAACTCAGCG  
TCCATCTGTTGATGTCATCTCTGGTTTTGATTAAGGCCAATGTTCCATCTCGTGTAGCATTTCGGGTTTC  
ATCAGGAACAGACTCCCGTACGATTTTGGATGAAAATGGAGCAGAAAAACTTCTTGGTCGAGGAGACAT  
GCTCTTTAAACCGATTGATGAAAATCATCCAGTTCGCTCCAAGGCTCCTTTATCTCGGATGACGATGT  
TGAGCGCATTTGGAACCTCATCAAGACTCAGGCAGATGCAGACTACGATGAGAGTTTTGATCCAGGTGA  
GGTTTTCTGAAAATGAAGGAGAATTTTCGGATGGAGATGCTGGTGGTGTATCCGCTTTTTGAAGAAGCTAA  
GTCTTTGGTTATCGAAACACAGAAAGCCAGTGCCTCTATGATTCAGCGTCGTTTATCAGTTGGATTAA  
CCGTGCGACCCGTCTCATGGAAGAACTGGAGATAGCAGGTGTCATCGGTCCAGCTGAAGGTACCAAACC  
TCGAAAAGTGTTACAACAA

**SP078 amino acid (SEQ ID NO:132)**

RGFAKWWEHERRKEERFVKQEEKARQKAEKEARLEQEETEKALLDLPPVDMETGEILTEEAVQNLPPI  
PEEKWVEPEIILPQAEKLFPEQEDSDDEDVQVDFSAKEALEYKLPQLFAPDKPKDQSEKKIVREN  
IKILEATFASFGIKVTERAEIGPSVTKYEVKPAVGVRVNRISNLSDDLALALAAKDVRIEAPIPGKSL  
IGIEVPNSDIATVSFRELWEQSQTKAENFLEIPLGKAVNGTARAFDLSKMPHLLVAGSTSGSKSVAVNG  
IIASILMKARPDQVKFMMVDPKMVELSVYNDIPHLIPVVTNPRKASKALQKVVDENRYELFAKVG  
RNIAGFNAKVVEFNSQSEYKQIPLPFIVVIVDELADLMMVASKEVEDAIIRLGQKARAAGIHMILATQR  
PSVDVISGLIKANVPSRVAFVSSGTDSTILDENGAEKLLGRGDMLEFKPIDENHPVRLQGSFISDDDV  
ERIVNFIKTQADADYDESFDPEVSENEGEFSDDGAGDPLFEEAKSLVIETQKASASMIQRRLSVGFN  
RATRLMEELEIAGVIGPAEGTKPRKVLQQ

**SP079 nucleotide (SEQ ID NO:133)**

TCAAAAAGAGAAGGAAAACCTTGGTTATTGCTGGGAAAATAGGTCCAGAACCAGAAATTTTGCCCAATAT  
GTATAAGTTGCTGATTGAAGAAAATACCAGCATGACTGCGACTGTAAACCGAATTTTGGGAAGACAAG  
CTTCCTTTTGAAGCTCTGAAAAAAGCGATATTGACATCTATCCTGAATTTACTGGTACGGTGACTGA  
AAGTTTGCTTCAACCATCACCCAAGGTGAGTCATGAACCAGAACAGGTTTATCAGGTGGCGCGTGATGG  
CATTGCTAAGCAGGATCATCTAGCCTATCTCAAACCCATGTCTTATCAAAACACCTATGCTGTAGCTGT  
TCCGAAAAGATTGCTCAAGAATATGGCTTGAAGACCATTTCAGACTTGAAAAAAGTGGAAGGCGAGTT  
GAAGGCAGGTTTTTACACTCGAGTTTAAACGACCGTGAAGATGGAAATAAGGGCTTGCAATCAATGTATGG  
TCTCAATCTCAATGTAGCGACCATTGAGCCAGCCCTTCGCTATCAGGCTATTTCAGTCAGGGGATATTCA  
AATCACGGATGCCTATTTCGACTGATGCGGAATTGGAGCGTTATGATTTACAGGTCTTGGAAGATGACAA  
GCAACTCTTCCCACCTTATCAAGGGGCTCCACTCATGAAAGAAGCTCTTCTCAAGAAACACCCAGAGTT  
GGAAAGAGTTCTTAATACATTGGCTGGTAAGATTACAGAAAGCCAGATGAGCCAGCTCAACTACCAAGT  
CGGTGTTGAAGGCAAGTCAGCAAAGCAAGTAGCCAAGGAGTTTCTCCAAGAACAAGGTTTGTGAAGAA  
A

**SP079 amino acid (SEQ ID NO:134)**

QKEKENLVIAGKIGPEPEILANMYKLLIEENTSMATVKNFNGKTSFLYEALKKGDIDIDYPEFTGTVTE  
SLLQSPKVSHEPEQVYQVARDGIKQDHLAYLKPMSYQNTYAVAVPKKIAQEYGLKTIISDLKKVEGQL  
KAGFTLEFNDREDGNKGLQSMYGLNLNVATIEPALRYQAIQSGDIQITDAYSTDAELERYDLQVLEDDK  
QLFPPYQGAPLMKEALLKKHPELERVNLNTLAGKITESQMSQLNLYQVGVGKSAKQVAKEFLQEQLLKK

**SP080 nucleotide (SEQ ID NO:135)**

ACGTTCTATTGAGGACCACTTTGATTCAAACCTTGAATTTGGAATATAACCTCAAAGAAAAAGGGAAAAC  
AGATCTTTTGAAGCTAGTTGATAAAACAACCTGACATGCGTCTGCATTTTATCCGCCAAACTCATCCACG  
CGGTCTCGGAGATGCTGTTTTGCAAGCCAAGGCTTTTCGTCGGAAATGAACCTTTTGTGCTTATGCTTGG  
TGATGACTTGATGGATATCACAGACGAAAAGGCTGTTCCACTTACCAAACAACCTCATGGATGACTACGA  
GCGTACCCACGCGTCTACTATCGCTGTCTATGCCAGTCCCTCATGACGAAGTATCTGCTTACGGGGTTAT  
TGCTCCGCAAGGCGAAGGAAAAGATGGTCTTTACAGTGTGAAACCTTTGTTGAAAAACAGCTCCAGA  
GGACGCTCCTAGCGACCTTGCTATTATCGGACGCTACCTCCTCAGCGCTGAAATTTTGTAGATTCTCGA  
AAAGCAAGCTCCAGGTGCAGGAAATGAAATTCAGCTGACAGATGCAATCGACACCCTCAATAAAACACA  
ACGTGTATTTGCTCGTGAGTTCAAAGGGGCTCGTTACGATGTCGGAGACAAGTTTGGCTTCATGAAAAC  
ATCCATCGACTACGCCCTCAAACACCCACAAGTCAAAGATGATTGAAGAATTACCTCATCCAACCTTGG  
AAAAGAATTGACTGAGAAGGAA

Table 1

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**SP080 amino acid (SEQ ID NO:136)**

RSIEDHFDNSNFELEYNLKEKGKTDLLKLVDKTTDMRLHFIRQTHPRGLGDAVLQAKAFVGNPEFVVMLG  
DDLMDITDEKAVPLTKQLMDDYERHASTIAVMPVPHDEVSAYGVIAPQGEKDGGLYSVETVFEKPAPE  
DAPSDLAIIGRYLLTPEIFEILEKQAPGAGNEIQLTDAIDTLNKTQRVFAREFKGARYDVGDKFGFMKT  
SIDYALKHPQVKDDLKNYLIQLGKELTEKE

**SP081 nucleotide (SEQ ID NO:137)**

CGCTCAAAATACCAGAGGTGTTTCAGCTAATCGAGCACGTTTCTCCTCAAATGTTGAAAGCCCAATTGGA  
GAGTGTCTTTTCTGATATTCCACCTCAGGCTGTAAAACTGGAATGTTGGCTACTACTGAAATCATGGA  
AATCATCCAACCCATCTTAAAAAACTGGATTGTCCCTATGTCCTTGATCCTGTTATGGTTGCTACAAG  
TGGAGATGCCTTGATTGACTCAAATGCTAGAGACTATCTCAAAACAACTTACTACCTCTAGCAACTAT  
TATTACGCCAAATCTTCCTGAAGCAGAAGAGATTGTTGGTTTTTCATCCATGACCCCGAAGACATGCA  
GCGTGCTGGTCGCTGATTTTAAAGAATTTGGTCCTCAGTCTGTGGTTATCAAAGCGGACATCTCAA  
AGGTGGTGCTAAAGATTTCCTCTTTACCAAGAATGAACAATTTGTCTGGGAAAGCCACGAATTCAAAC  
CTGTACACCCATGGTACT

**SP081 amino acid (SEQ ID NO:138)**

AQNTRGVQLIEHVSPQMLKAQLESVFS DIPPAVKTGMLATTEIMEIIQPYLKKLDCPYVLDPMVMVATS  
GDALIDS NARDYLKTNLLPLATIIITPNLPEAEIEIVGFSIHDPEDMQRAGRLILKEFGPQSVVIKGGHLK  
GGAKDFLFTKNEQFVWESPRIQTCHTHGT

**SP082 nucleotide (SEQ ID NO:139)**

AATTGTACAATTAGAAAAAGATAGCAAATCAGACAAAGAACAAGTTGATAAACTATTTGAATCATTTGA  
TGCATCTTCAGATGAATCTATTTCTAAATTAAGAAGCTATCTGAACTTCACTTAAACCGATGCAGG  
TAAAGACTATCTTAATAACAAAGTCAAAGAATCATCTAAAGCAATTGTAGATTTTCATTTGCAAAAAGG  
TTTGGCTTATGATGTTAAAGATTGAGATGACAAATTTAAAGATAAAGCAACTCTTGAAACAAATGTAAA  
AGAAATTACAAAACAAATTGATTTTATCAAAAAAGTTGATGAACTTTTAAACAAGAGAATTTGGAAGA  
AACTCTTAAATCTCTAAATGATCTTGTGATAAATATCAAAAACAAATCGAACTTTTGAAGAAAGAAGA  
AGAAAAAGCTGCTGAAAAAGCTGCTGAAAAAGCAAAGGAATCTTCTAGTCAAAGTAATTTCTTGGTAG  
TGCTTCTAATGAGTCTTATAATGGATCTTCCAATTCAAATGTAGATTATAGTTTCTGAACAACTAA  
TGGATATTCAAATAATTATGGCGGTCAAGATTATCTGGTTCAGGAGATAGTTCAACAAATGGTGGATC  
ATCAGAACAAATATTCATCTAGCAATTCAAACAGCGGAGCAAATAATGTCTACAGATATAAAGGCACTGG  
TGCTGACGGCTATCAAAGATACTACTACAAAGATCATAATAATGGAGATGTGTATGATGACGATGGAAA  
TTACCTTGGGAACCTTTGGTGGCGGCATTGCAGAACCTAGTCAACGC

**SP082 amino acid (SEQ ID NO:140)**

IVQLEKDSKSDKEQVDKLFESFDASSDESISKLELSETSLKTDAGKDYLNKNVKESSKAIVDFHLQKG  
LAYDVKDSDDKFKDKATLETNVKEITKQIDFIKKVDETFKQENLEETLKS LNDLVDKYQKQIELLKKEE  
EKAAEKAAEKAKESSQSNSSGSASNESYNGSSNSNVDSSEQTNGYSNNYGGQDYSGSGDSSTNGGS  
SEQYSSSNSNSGANNVRYRYKGTGADGYQRYYYKDHNNGDVYDDDGNYLGNFGGGIAEPSQR

**SP083 nucleotide (SEQ ID NO:141)**

TCTGACCAAGCAAAAAGAAGCAGTCAATGACAAAGGAAAAGCAGCTGTTGTTAAGGTGGTGGAAAGCCA  
GGCAGAACTTTATAGCTTAGAAAAGAATGAAGATGCTAGCCTAAGAAAAGTTACAAGCAGATGGACGCAT  
CACGGAAGAACAGGCTAAAGCTTATAAAGAATACAAATGATAAAAAATGGAGGAGCAAATCGTAAAGTCAA  
TGAT

**SP083 amino acid (SEQ ID NO:142)**

LTKQKEAVNDKGKAAVVKVVESQAELYSLEKNEDASLRKLQADGRITEEQAKAYKEYNDKNGGANRKVN  
D

**SP084 nucleotide (SEQ ID NO:143)**

GTCCGGCTCTGTCCAGTCCACTTTTTCAGCGGTAGAGGAACAGATTTTCTTTATGGAGTTTGAAGAACT  
CTATCGGGAAACCCAAAAACGCAGTGTAGCCAGTCAGCAAAAAGACTAGTCTGAACCTTAGATGGGCAGAC  
GCTTAGCAATGGCAGTCAAAAGTTGCCAGTCCCTAAAGGAATTCAGGCCCATCAGGCCAAAGTATTAC  
ATTTGACCGAGCTGGGGGCAATTCGTCCCTGGCTAAGGTTGAATTTAGACCAGTAAAGGAGCGATTTCG  
CTATCAATTATATCTAGGAAATGGAAAAATTAACGCATTAAGGAAACAAAAAAT

Table 1

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**SP084 amino acid (SEQ ID NO:144)**

SGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQOKTSLNLDGQTLNNGSQKLPVPKGIQAPSGQSIT  
FDRAGGNSSLAKVEFQTSKGAIRYQLYLGNNGKIKRIKETKN

**SP085 nucleotide (SEQ ID NO:145)**

GGGACAAATTCAAAAAAATAGGCAAGAGGAAGCAAAAATCTTGCAAAAGGAAGAAGTCTTGAGGGTAGC  
TAAGATGGCCCTGCAGACGGGGCAAAATCAGGTAAGCATCAACGGAGTTGAGATTCAGGTATTTCTAG  
TGAAAAAGGATTGGAGGTCTACCATGGTTCAGAACAGTTGTTGGCAATCAAAGAGCCA

**SP085 amino acid (SEQ ID NO:146)**

GQIQKNRQEEAKILQKEEVLRVAKMALQTGQNVQSINGVEIQVFSSEKGLEVYHGSEQLLAIKEP

**SP086 nucleotide (SEQ ID NO:147)**

TCGCTACCAGCAACAAAGCGAGCAAAAGGAGTGGCTCTTGTGTTGTGGACCAACTTGAGGTAGAATTAGA  
CCGTTTCGCAGTTTCGAAAAAGTAGAAGGCAATCGCCTATACATGAAGCAAGATGGCAAGGACATCGCCAT  
CGGTAAGTCAAAGTCAGATGATTTCCGTAAAACGAATGCTCGTGGTTCGAGGTATCAGCCTATGGTTTA  
TGGACTCAAATCTGTACGGATTACAGAGGACAATCAACTGGTTCGCTTTCATTTCCAGTTCCAAAAGG  
CTTAGAAAGGGAGTTCATCTATCGTGTGAAAAAGAAAAAGT

**SP086 amino acid (SEQ ID NO:148)**

RYQQQSEQKEWLLFVDQLEVELDRSQFEKVEGNRLYMKQDGKDIAIGKSKSDDFRKTNARGRGYQPMVY  
GLKSVRITEDNQLVRFHFQFQKGLEREFYRVEKEKS

**SP087 nucleotide (SEQ ID NO:149)**

GAACCGACAAGTCGCCCCTATCAAGACTATGCTTTGAATAAAGAAAAATTGGTTGCTTTTGCTATGGC  
TAAACGAACCAAAGATAAGGTTGAGCAAGAAAGTGGGGAACAGTTTTTAATCTAGGTCAGGTAAGCTA  
TCAAAACAAGAAAAGTGGCTTAGTGACGAGGGTTCGTACCGATAAGAGCCAATATGAGTTTCTGTTTCC  
TTCAGTCAAATCAAAGAAGAGAAAAGAGATAAAAAGGAAGAGGTAGCGACCGATTCAAGCGAAAAAGT  
GGAGAAGAAAAATCAGAAGAGAAGCCTGAAAAGAAAGAGAATTCA

**SP087 amino acid (SEQ ID NO:150)**

NRQVAHYQDYALNKEKLVAFAMAKRTKDKVEQESGEQFFNLGQVSYQNKKTGLVTRVRTDKSQYEFLFP  
SVKIKEEKRDKEEVATDSSEKVEKKKSEEKPEKKENS

**SP088 nucleotide (SEQ ID NO:151)**

GGTTGTGGCTGGCAATATATCCCGTTTCCATCTAAAGGTAGTACAATTGGTCCTTACCCAAATGGTAT  
CAGATTAGAAGTTTTCCAAAGTCAGAGTGGTACTACTTTCGATAAAAATGGAGTGCTACAAGAGTTTGT  
TGGTTGGAACATTAGAGATTAAACTAAAGACAGTGTGGAAGAAAGTACGGGGAACCGTGAAGA  
TTCAGAAGATAAAGAAGAGAAGCGTTATTATACGAACTATTACTTTAATCAAATCATTTCTTTAGAGAC  
AGTTTGGCTTTATGATGACTCTAAGTATTATCTAGCTAAGACGGAAATTAATGGAGAAAACCTACCT  
TGGTGGTGAAGACGTCGCGGGTGGATAAACGATGATTCGACTTGGTACTACCTAGATCCAACAACTGG  
TATTATGCAACAGGTTGGCAATATCTAGGTAATAAGTGGTACTACCTCCGTTCCCTCAGGAGCAATGGC  
CACTGGCTGGTATCAGGAAGGTACCCTTGGTATTATTTAGACCACCCAAATGGCGATATGAAAACAGG  
TTGGCAAAACCTTGGGAACAAATGGTACTATCTCCGTTTCATCAGGAGCTATGGCAACTGGTTGGTATCA  
AGATGGTTCAACTTGGTACTACCTAAATGCAGGTAATGGAGACATGAAGACAGGTGGTTCCAGGTCAA  
TGGCAACTGGTACTATGCTTATAGCTCAGGTGCTTTGGCAGTGAATACGACCGTAGATGGCTATTCTGT  
CAACTATAATGGCGAATGGGTTCCG

**SP088 amino acid (SEQ ID NO:152)**

VVGWQYIPFPKSGSTIGPYPNGIRLEGFPKSEWYYFDKNGVLQEFVGVWKTLEIKTKDSVGRKYGEKRED  
SEDKEEKRYYTNYFNQNHSLGTWLYDQSNWYYLAKTEINGENYLGGERAGWINDDSTWYYLDPTTG  
IMQTGWQYLGKWWYLRSSGAMATGWYQEGTTWYYLDHPNGDMKTGWQNLGNKWWYLRSSGAMATGWYQ  
DGSTWYYLNAGNGDMKTGWYFQVNGNWYYAYSSGALAVNTTVDGYSVNYNGEWR

**SP089 nucleotide (SEQ ID NO:153)**

GGCCAAATCAGAATGGGTAGAAGACAAGGGAGCCTTTTATTATCTTGACCAAGATGGAAAGATGAAAAG  
AAATGCTTGGGTAGGAACCTTCCTATGTTGGTGAACAGGTGCCAAAGTAATAGAAGACTGGGTCTATGA  
TTCTCAATACGATGCTTGGTTTTATATCAAAGCAGATGGACAGCACGCAGAGAAAAGAAATGGCTCCAAAT

Table 1

TAAAGGGAAGGACTATTATTTCAAATCCGGTGGTTATCTACTGACAAGTCAGTGGATTAATCAAGCTTA  
TGTGAATGCTAGTGGTGCCAAAGTACAGCAAGGTTGGCTTTTTGACAAACAATACCAATCTTGGTTTTA  
CATCAAAGAAAATGGAACTATGCTGATAAAGAATGGATTTTCGAGAATGGTCACTATTATTATCTAAA  
ATCCGGTGGCTACATGGCAGCCAATGAATGGATTGGGATAAGGAATCTTGGTTTTATCTCAAATTTGA  
TGGGAAAATGGCTGAAAAAGAATGGGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCCGG  
TGGTTACATGACAGCCAATGAATGGATTGGGATAAGGAATCTTGGTTTTATCTCAAATCTGATGGGAA  
AATAGCTGAAAAAGAATGGGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCCGGTGGTTA  
CATGACAGCCAATGAATGGATTGGGATAAGGAATCTTGGTTTTACCTCAAATCTGATGGGAAAATAGC  
TGAAAAAGAATGGGTCTACGATTCTCATAGTCAAGCTTGGTACTACTTCAAATCTGGTGGCTACATGGC  
GAAAAATGAGACAGTAGATGGTTATCAGCTTGGGAAGCGATGGTAAATGGCTTGGAGGAAAAACTACAAA  
TGAAAATGCTGCTTACTATCAAGTAGTGCCTGTTACAGCCAATGTTTATGATTGAGATGGTGAAAAGCT  
TTCTTATATATCGCAAGGTAGTGTCTGATGGCTAGATAAGGATAGAAAAAGTGATGACAAGCGCTTGGC  
TATTACTATTTCTGGTTTGTGAGGCTATATGAAAACAGAAGATTACAAGCGCTAGATGCTAGTAAGGA  
CTTTATCCCTTATTATGAGAGTGATGGCCACCGTTTTTATCACTATGTGGCTCAGAATGCTAGTATCCC  
AGTAGCTTCTCATCTTTCTGATATGGAAGTAGGCAAGAAATATTATTCGGCAGATGGCCTGCATTTTGA  
TGGTTTTAAGCTTGAGAATCCCTTTCCTTTTCAAAGATTTAACAGAGGCTACAACTACAGTGCTGAAGA  
ATTGGATAAGGTATTTAGTTTGTCTAAACATTAAACATAGCCTTTTGGAGAACAAGGGCGCTACTTTTAA  
GGAAGCCGAAGAACATTACCATATCAATGCTCTTTATCTCCTTGGCCATAGTGCCCTAGAAAGTAAGTG  
GGGAAGAAGTAAAATTGCCAAAGATAAGAATAATTTCTTTGGCATTACAGCCTATGATACGACCCCTTA  
CCTTTCTGCTAAGACATTGATGATGTGGATAAGGGAATTTAGGTGCAACCAAGTGGATTAAGGAAAA  
TTATATCGATAGGGGAAGAACTTTCTTGGAAACAAGGCTTCTGGTATGAATGTGGAATATGCTTCAGA  
CCCTTATTGGGGCGAAAAAATTGCTAGTGTGATGATGAAAATCAATGAGAAG

## SP089 amino acid (SEQ ID NO:154)

AKSEWVEDKGAFYYLDQDGKMKRNWVGTSYVGATGAKVIEDWVYDSQYDAWFYIKADGQHAKEKWLQI  
KGKDYYFKSGGYLLTSQWINQAYVNASGAKVQQGWLFQKQYQSWFYIKENGNYADKEWIFENGHYYYLK  
SGGYMAANEWIWDKESWFLKFDGKMAEKWVYDSHSQAWYYFKSGGYMTANEWIWDKESWFLKSDGK  
IAEKWVYDSHSQAWYYFKSGGYMTANEWIWDKESWFLKSDGKIAEKWVYDSHSQAWYYFKSGGYMA  
KNETVDGYQLGSDGKWLGGKTTNENAAYYQVVPVTANVYDSGGEKLSYISQGSVVWLDKDRKSDDKRLA  
ITISGLSGYMKTEDLQALDASKDFIPYYESDGHFRFYHYVAQNASIPVASHLSDMEVGKKYYSADGLHFD  
GFKLENPFLFKDLTEATNYSAEELDKVFSLLNINNSLLENKGATFKEAEEHYHINALYLLAHSALSNW  
GRSKIADKNNFFGITAYDTTPYLSAKTFDDVDKGILGATKWIKENYIDRGRTFLGNKASGMNVEYASD  
PYWGEKIASVMMKINEK

## SP090 nucleotide (SEQ ID NO:155)

ATTTGCAGATGATTCTGAAGGATGGCAGTTTGTCCAAGAAAATGGTAGAACCTACTACAAAAGGGGGA  
TCTAAAAGAAACCTACTGGAGAGTGATAGATGGGAAGTACTATTATTTTGATCCTTTATCCGGAGAGAT  
GGTTGTGCGCTGGCAATATATACCTGCTCCACACAAGGGGGTTACGATTGGTCCTTCTCCAAGAATAGA  
GATTGCTCTTAGACCAGATTGGTTTTATTTTGGTCAAGATGGTGTATTACAAGAATTTGTTGGCAAGCA  
AGTTTTTAGAAGCAAAAACCTGCTACGAATACCAACAAACATCATGGGGAAGAATATGATAGCCAAGCAGA  
GAAACGAGTCTATTATTTTGAAGATCAGCGTAGTTATCATACTTTAAAACTGGTTGGATTATGAAGA  
GGGTCAATTGGTATTATTTACAGAAGGATGGTGGCTTTGATTTCGCGCATCAACAGATTGACGGTTGGAGA  
GCTAGCACGTGGTTGGGTTAAGGATTACCCTCTTACGTATGATGAAGAGAAGCTAAAAGCAGCTCCATG  
GTACTATCTAAATCCAGCAACTGGCATTATGCAAAACAGGTTGGCAATATCTAGGTAATAGATGGTACTA  
CCTCCATTCGTCAGGAGCTATGGCAACTGGCTGGTATAAGGAAGGCTCAACTTGGTACTATCTAGATGC  
TGAAAATGGTGATATGAGAACTGGCTGGCAAAACCTTGGGAACAAATGGTACTATCTCCGTTTCATCAGG  
AGCTATGGCAACTGGTTGGTATCAGGAAAGTTCGACTTGGTACTATCTAAATGCAAGTAATGGAGATAT  
GAAAACAGGCTGGTTCCAAGTCAATGGTAACTGGTACTATGCCTATGATTGAGGTGCTTTAGCTGTAA  
TACCACAGTAGGTGGTTACTACTTAAACTATAATGGTGAATGGGTAAAG

## SP090 amino acid (SEQ ID NO:156)

VFADDSEGWFVQENGRTYYKKGDLKETYWRVIDGKYYYFDPLSGEMVVGWQYIPAPHKGVITIGPSPRI  
EIALRPDWFYFGQDGVLFQEFVGKQVLEAKTATNTNKKHHGEEYDSQAEKRVYFEDQRSYHTLKTGWIYE  
EGHWYYLQKDGGFDSRINRLTVGELARGWVKDYPLTYDEEKLKAAPWYYLNPATGIMQTGWQYLGNRWY  
YLHSSGAMATGWYKEGSTWYYLDAENGDMRTGWQNLGNKWYYLRSSGAMATGWYQESSTWYYLNASNGD  
MKTGWFFQVNGNWYYAYDSGALAVNTTVGGYYLNYNGEWVK

Table 1

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**SP091 nucleotide (SEQ ID NO:157)**

TGTCGCTGCAAATGAAACTGAAGTAGCAAAAACCTTCGCAGGATACAACGACAGCTTCAAGTAGTTCAGAG  
GCAAAATCAGTCTTCTAATAAAACGCAAACGAGCGCAGAAGTACAGACTAATGCTGCTGCCCACTGGGA  
TGGGGATTATTATGTAAAGGATGATGGTTCCTAAAGCTCAAAGTGAATGGATTTTGGACAACTACTATAA  
GGCTTGGTTTTATATTAATTCAGATGGTCGTTACTCGCAGAATGAATGGCATGGAAATTACTACCTGAA  
ATCAGGTGGATATATGGCCCAAACGAGTGGATCTATGACAGTAATTACAAGAGTTGGTTTTATCTCAA  
GTCAGATGGGGCTTATGCTCATCAAGAATGGCAATTGATTGGAAATAAGTGGTACTACTTCAAGAAGTG  
GGTTTACATGGCTAAAAGCCAATGGCAAGGAAGTTATTTCTTGAATGGTCAAGGAGCTATGATGCAAAA  
TGAATGGCTSCTATGATCCAGCCTATTCTGCTTATTTTATCTAAAATCCGATGGAACTTATGCTAACC  
AAGAGTGGCAAAAAGTGGGCGGCAAAATGGTACTATTCTCAAGAAGTGGGGCTATATGGCTCGGAATGAGT  
GGCAAGGCAACTACTATTTGACTGGAAGTGGTGGCCTGGCGACTGACGAAGTGATTATGGATGGTACTC  
GCTATATCTTTGCGGCTCTGGTGAGCTCAAAGAAAAAAGATTTGAATGTCGGCTGGGTTTCACAGAG  
ATGGTAAGCGCTATTTCTTTAATAATAGAGAAGAACAAGTGGGAACCGAACATGCTAAGAAAAGTCATTG  
ATATTAGTGAGCACAATGGTCGTATCAATGATTGGAAAAAGGTTATTGATGAGAACGAAGTGGATGGTG  
TCATTGTTCTGCTAGGTTATAGCGGTAAAGAAGACAAGGAATTGGCGCATAACATTAAGGAGTTAAACC  
GTCTGGGAATTCCTTATGGTGTCTATCTCTATACCTATGCTGAAAATGAGACCGATGCTGAGAGTGACG  
CTAAACAGACCATTGAACCTTATAAAGAAATACAATATGAACCTGTCTTACCCTATCTATTATGATGTTG  
AGAATTGGGAATATGTAAATAAGAGCAAGAGAGCTCCAAGTGATACAGGCACTTGGGTAAAATCATCA  
ACAAGTACATGGACACGATGAAGCAGGCGGGTTATCAAAATGTGTATGTCTATAGCTATCGTAGTTTAT  
TACAGACGCGTTTAAAAACCCAGATATTTTAAACATGTAAACTGGGTAGCGGCCCTATACGAATGCTT  
TAGAATGGGAAAACCTCATTATTCAGGAAAAAAGGTTGGCAATATACCTCTTCTGAATACATGAAAG  
GAATCCAAGGGCGCTAGATGTCAGCGTTTGGTAT

**SP091 amino acid (SEQ ID NO:158)**

VAANETEVAKTSQDTTASSSSEQNQSSNKTQTSAEVQTNAAAHWDGDYVYKDDGSKAQSEWIFDNYYK  
AWFYINSDGRYSQNEWHGNYLKSQGYMAQNEWIYDSNYKSWFYLKSDGAYAHQEWQLIGNKWWYFKKW  
GYMAKSQWQGSYFLNQGAMMONEWLYDPAYSAYFYLKSDGTYANQEWQKVGGKWYFKKWGYMARNEW  
QGNYYLTGSGAMATDEVIMDGTTRYIFAASGELKEKKDLNVGWVHRDQKRYFFNNREEQVGEHAKKVID  
ISEHNGRINDWKVIDENEVDGVIIVRLGYSGKEDKELAHNIKELNRLGIPYGVYLYTYAENETDAESDA  
KQTIELIKKYNMNLSPYIYDVENWEYVNKSKRAPSDTGTWVKIINKYMDTMKQAGYQNVVYSYRSL  
QTRLKHPDILKHVNWVAAYTNALEWENPHYSKKGWQYTSSEYMKGIQGRVDVSVWY

**SP092 nucleotide (SEQ ID NO:159)**

TACGTCTCAGCCTACTTTTGTAAGAGCAGAAGAATCTCCACAAGTTGTGCAAAAATCTTCATTAGAGAA  
GAAATATGAGGAAGCAAAAGCAAAAGCTGATACTGCCAAGAAAGATTACGAAACGGCTAAAAAGAAAGC  
AGAAGACGCTCAGAAAAAGTATGAAGATGATCAGAAGAGAACTGAGGAGAAAGCTCGAAAAAGACAGA  
AGCATCTCAAAAATTGAATGATGTGGCGCTTGTTGTTCAAAATGCATATAAAGAGTACCGAGAAGTTCA  
AAATCAACGCTAGTAAATATAAATCTGACGCTGAATATCAGAAAAAATTAACAGAGGTCGACTCTAAAT  
AGAGAAGGCTAGGAAAGACGCAACAGGACTTGCAAAATAAATTTAATGAAGTAAGAGCAGTTGTAGTTCC  
TGAACCAAATGCGTTGGCTGAGACTAAGAAAAAGCAGAAGAAGCTAAAGCAGAAGAAAAAGTAGCTAA  
GAGAAAAATATGATTATGCAACTCTAAAGGTAGCACTAGCGAAGAAAGAAGTAGAGGCTAAGGAACTTGA  
AATTGAAAAACTTCAATATGAAATTTCTACTTTGGAACAAGAAGTTGCTACTGCTCAACATCAAGTAGA  
TAATTTGAAAAAACTTCTTGCTGGTGCGGATCCTGATGATGGCACAGAAGTTATAGAAGCTAAATTTAA  
AAAAGGAGAAGCTGAGCTAAACGCTAAACAAGCTGAGTTAGCAAAAAAACAACAGAACTTGA AAAACT  
TCTTGACAGCCTTGATCCTGAAGGTAAGACTCAGGATGAATTAGATAAAGAAGCAGAAGAAGCTGAGTT  
GGATAAAAAAGCTGATGAACCTTCAAAATAAAGTTGCTGATTTAGAAAAAGAAATTAGTAACCTTGAAAT  
ATTACTTGGAGGGGCTGATNCTGAAGATGATACTGCTGCTCTTCAAAATAAATTAGCTACTAAAAAAGC  
TGAATTGGAAAAAAGCTCAAAAAGAATTAGATGCAGCTCTTAATGAGTTAGGCCCTGATGGAGATGAAGA  
AGAAACTCCAGCGCCGGCTCCTCAACCAGAGCAACCAGCTCCTGCACCAAAACCAGAGCAACCAGCTCC  
AGCTCCAAAACCAGAGCAACCAGCTCCTGCACCAAAACCAGAGCAACCAGCTCCAGCTCCAAAACCAGA  
GCAACCAGCTCCAGCTCCAAAACCAGAGCAACCAGCTAAGCCGAGAGAAACCAGCTGAAGAGCCTACTCA  
ACCAGAAAAACCAGCCTCCAAAACAGGCTGGAACAAGAAAAACGGTATGTGGTATTTCTACAATAC  
TGATGGTTCAATGGCAATAGGTTGGCTCCAAAACACGGTTTCATGGTACTACCTAAACGCTAACGGCGC  
TATGGCAACAGGTTGGGTGAAAGATGGAGATACCTGGTACTATCTTGAAGCATCAGGTGCTATGAAAGC  
AAGCCAATGGTTCAAAGTATCAGATAAATGGTACTATGTCAACAGCAATGGCGCTATGGCGACAGGCTG  
GCTCCAATACAATGGCTCATGGTACTACCTCAACGCTAATGGTGATATGGCGACAGGATGGCTCCAATA  
CAACGGTTTCATGGTATTACCTCAACGCTAATGGTGATATGGCGACAGGATGGGCTAAAGTCAACGGTTC  
ATGGTACTACCTAAACGCTAACGGTGTATGGCTACAGGTTGGGCTAAAGTCAACGGTTCATGGTACTA

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SP092 amino acid (SEQ ID NO:160)

P093\_nucleotide (SEQ ID NO:161)

SP093 amino acid (SEQ ID NO:162)

SP094 nucleotide (SEQ ID NO:163)

SP094 amino acid (SEQ ID NO:164)

SP095 nucleotide (SEQ ID NO:165)

TAGGTCATATGGGACTTTTTTCTACACAAAAATAGGCTCCATAATATCTATAAGGGATTACCCACTA  
CAAATATTATAGAGCCGAAAATTACATCTAATATATGCAGACTACTTTGAAATGAAATTAATAAAATTT  
ATTAAAGGATGACACAAAAGTTTTTGAAAAATCTACATTCAAATTTGTAGAAGGATATAAAATATACCT

Table 1

GACAGAATCTAAAGAATCTGGAATTAAACAAATGGACAATGTCATAAAATATTTTGAAGTTTATTGAATC  
TAAAAGTATTGCTTTATATTTTCAAAAACGATTAAATGAGCTGATAGAT

**SP095 amino acid (SEQ ID NO:166)**

RSYGTFFLQQNRLHNIYKGFTHYKYRAENSHLIYADYFEMKLLKLLKDDTKVFEKSTFKFVEGYKIYL  
TESKESGIKQMDNVIKYFEFIESKSIALYFQKRLNELID

**SP096 nucleotide (SEQ ID NO:167)**

CAACGTTGAGAATTATTTGCGAATGTGTTTGATAGCATTGAGAATCAGACGTATCAAAATTTTGAGTG  
TTTATTAATCAATGATGGCTCTCCAGATCATTCACAAAATATGTGAAGAATTTGTAGAGAAAGATTC  
TCGTTTCAAATATTTTGAGAAAGCAAACGGCGGCTTTTCATCAGCTCGTAACCTAGGTATTGAATGTTT  
GGGGGGGGCGTACATTACTTTTGTAGACTC

**SP096 amino acid (SEQ ID NO:168)**

NVENYLRMCLDSIQNQTYQNFECCLINDGSPDHSSKICEEFVEKDSRFKYFEKANGGLSSARNLGIECS  
GGGVHYFCRL

**SP097 nucleotide (SEQ ID NO:169)**

CTACTATCAATCAAGTTCTTCAGCCATTGAGGCCACCATTGAGGGCAACAGCCAAACGACCATCAGCCA  
GACTAGCCACTTTTATTCAGTCTTATATCAAAAACTAGAAACCACCTCGACTGGTTTGACCCAGCAGAC  
GGATGTTCTGGCCTATGCTGAGAATCCAGTCAAGACAAGGTCGAGGGAATCCGAGATTTGTTTTTGAC  
CATCTTGAAGTCAGATAAGGACTTGAAAACCTGTTGTGCTGGTGACCAAATCTGGTCAGGTCATTCTAC  
AGATGACAGTGTGCAGATGAAAACCTTCTCTGATATGATGGCTGAGGATTGGTACCAAAGGCCATTCA  
TCAGGGAGCTATGCCTGTTTTGACTCCAGCTCGTAAATCAGATAGTCAGTGGGTCAATTCTGTCACTCA  
AGAACTTGTGATGCAAAGGGAGCCAATCTTGGTGTGCTTCGTTTGGATATTTCTTATGAAAACCTCTGGA  
AGCCTATCTCAATCAACTCCAGTTGGGGCAGCAGGGCTTTGCCTTCATTATCAATGAAAACCATGAATT  
TGCTTACCATCCTCAACACACAGTTTATAGTTTCGTCTAGCAAATGGAGGCTATGAAACCTTACATCGA  
TACAGGTCAGGGTTTACTCCTGGTCACAAATCCTACGTCAGTCAAGAGAAGATTGCAGGAACTGATTG  
GACGGTGCCTTGGCGTGTCAATTCGAAAAGTTAGACCAGGTTCCGGAGTCAG

**SP097 amino acid (SEQ ID NO:170)**

YYQSSSSAIEATIEGNSQTTISQTSHFIIQSYIKKLETTSTGLTQOTDVLAYAENPSQDKVEGIRDLEFLT  
ILKSDKDLKTVVLVTKSGQVISTDDSVQMKTSDDMMAEDWYQKAIHQGAMPVLT PARKSDSQWVISVTQ  
ELVDAGANLGVRLDISYETLEAYLNQLQLGQQGFAFIINENHEFVYHPQHTVYSSSSKMEAMKPYID  
TGQGYTPGHKSIVSQEKIAGTDWTVLGVSSLEKLDQVRSQ

**SP098 nucleotide (SEQ ID NO:171)**

GACAAAAACATTAAAACGTCCTGAGGTTTTATCACCTGCAGGGACTTTAGAGAAGCTAAAGGTAGCTGT  
TCAGTATGGAGCAGATGCTGTCTTTATCGGTGGTCAGGCCTATGGTCTTCGTAGCCGTGCGGGAAACTT  
TACTTTCGAACAGATGGAAGAAGGCGTGCAGTTTGC GGCCAAGTATGGTGCCAAGGTCTATGTAGCGGC  
TAATATGGTTATGCACGAAGGAAATGAAGCTGGTGTGCTGGTGAGTGGTTCCGTAAACTGCGTGATATCGG  
GATTGCAGCAGTTATCGTATCTGACCCAGCCTTGATTATGATTGCAGTGACTGAAGCACCAGGCCTTGA  
AATCCACCTTTCTACCCAAGCCAGTGCCACTAATGAAACCCCTTGAGTTCTGGAAAGAGCTAGGCTT  
GACTCGTGTGCTTTTAGCGCGTGAGGTTTCAATGGAAGAATTAGCTGAGATCCGCAAACGTACAGATGT  
TGAAATTGAAGCCTTTGTCCATGGAGCTATGTGTATTTTCACTCTGGACGTTGTACTCTTTCAAACCA  
CATGAGTATGCGTGATGCCAACCGTGGTGGATGTTCTCAGTCATGCCGTTGGAAATACGACCTTTACGA  
TATGCCATTTGGGAAAGAACGTAAGAGTTTGCAGGGTGAGATTCCAGAAGAATTTTCAATGTCAGCCGT  
TGACATGTCTATGATTGACCANATTCCAGATATGATTGAAAATGGTGTGGACAGTCTAAAAATCGAAGG  
ACGTATGNAGTCTATTTCACTANGTATCAACAGTAACCAACTGCTACAAGGCGGCTGTGGATGCCTATCT  
TGAAAGTCCTGAAAAGTTTGAAGCTATCAAACAAGACTTGGTGGACGAGATGTGGAAGGTTGCCAACG  
TGAAGTGGCTACAGGATTTTACTATGGTACACCATCTGAAAATGAGCAGTTGTTTGGTGTCTGCTGTAA  
AATCCCTGAGTACAAGTTTGTGCTGAAGTGGTTTCTTATGATGATGCGGCACAAACAGCAACTATTCG  
TCAACGAAACGTCATTAACGAAGGGGACCAAGTTGAGTTTATGGTCCAGGTTTCCGTCATTTTGAAC  
CTATATTGAAGATTTGCATGATGCTAAAGGCAATAAAATCGACCGCGCTCCAAATCCAATGGAAGTATT  
GACTATTAAAGTCCCACAACCTGTTCAATCAGGAGACATGGTTTCGAGCTCTTAAAGAGGGGCTTATCAA  
TCTTTATAAGGAAGATGGAACCAGCGTCACAGTTCGTGCT

Table 1

**SP098 amino acid (SEQ ID NO:172)**

TKTLKRPEVLSFAGTLEKLKVAVQYGADAVFIGGQAYGLRSRAGNFTFEQMEEGVQFAAKYGAKVYVAA  
NMVMHEGNEAGAGEWFRKLRLDIGIAAVIVSDPALIMIAVTEAPGLEIHLSTQASATNYETLEFWKELGL  
TRVVLAREVSMEEELAEIRKRTDVEIEAFVHGAMCISYSGRCTLSNHMSMRDANRGGCSQSCRWKYDLYD  
MPFGKERKSLQGEIPEEFSMSAVDMSMIDXI PDMIENGVDLSKIEGRMXSIHXVSTVTNCKYAAVDAYL  
ESPEKFEAIKQDLVDEMVKVAQRELATGFYFGTPESENEQLFGARRKIPEYKFVAEVVSYDDAAQTATIR  
QRNVINEGDQVEFYGPGRHFETIYIEDLHDAKGNKIDRAPNPMELLTIKVPQPVQSGDMVRALKEGLIN  
LYKEDGTSVTVRA

**SP099 nucleotide (SEQ ID NO:173)**

TTCTCAGGAGACCTTTAAAAATATCACCAATAGCTTCTCCATGCAAATCAATCGTCGCGTCAACCAAGG  
AACGCCTCGTGGTCTGCGGAATATCAAGGGTGAAGACATCAAAAAATCACCGAAAACAAGGCCATTGA  
GTCTTATGTCAAACGTATCAACGCTATCGGAGATTTGACTGGATATGACCTGATTGAAACGCCAGAAAC  
CAAGAAGAATCTCACTGCTGATCGTGCCAAGCGTTTTGGAAGTAGCTTGATGATTACAGGTGTCAATGA  
CTCCTCTAAAGAAGACAAGTTTGTCTCTGGTTCTTATAAACTAGTCGAAGGAGAGCACTTAACCAACGA  
CGACAAGGATAAAATCCTCTTGCACAAGGACTTGGCAGCCAAACACGGCTGGAAAGTAGGGGACAAGGT  
TAACTGGACTCTAATATCTACGATGCAGATAATGAAAAAGGAGCCAAGGAAACAGTTGAAGTGACAAT  
CAAGGGACTCTTTGATGGTCATAATAAGTCAGCAGTAACCTACTCACAAGAACTTTACGAAAACACAGC  
TATTACAGACATTCACACTGCTGCAAACTTTATGGATACACAGAAGACACAGCCATTTATGGGGACGC  
AACCTTCTTTGTAACAGCAGACAAGAACTTGGATGATGTTATGAAAGAGTTGAATGGCATCAGTGGTAT  
CAACTGGAAGAGCTACACACTCGTCAAGAGCTCCTCTAACTACCCAGCTCTTGAGCAATCTATCTCTGG  
TATGTACAAGATGGCCAAAC

**SP099 amino acid (SEQ ID NO:174)**

SQETFKNITNSFSMQINRRVNQGTPRGAGNIKGEDIKKITENKAIESYVKRINAIGDLTGydLIETPET  
KKNLTADRAKRFGSSLMITGVNDSSKEDKFVSGSYKLVEGEHLTNDKDKILLHKDLAAKHGKVGDKV  
KLDSNIYDADNEKGAKETVEVTIKGLFDGHNKSAVTYSQELYENTAITDIHTAAKLYGYTEDTAIYGDA  
TFFVTADKNLDDVMKELNGISGINWKS YTLVKSSSNYPALQSI SGM YKMAN

**SP100 nucleotide (SEQ ID NO:175)**

AGTAAATGCGCAATCAAATTCATTAATATTAATAGATGAACCTGAAATCTCACTTCATCCGAGTGCAAT  
CTATAAATTTAAAGAGTTTTTACTTCAAGAGTGTTTAAATAAAAAACATCAAATTATTATCACTACACA  
TTCTACACAACCTATAAAAGATTTTCTTAGAGAAGCCGTGAACTTTTAGTGAAAAACGGAGAAAAGGT  
AGATGTTATTGAAAATATTGATTATCAGGATGCATTTTTTGAATTAGGTGATGTGTATCATTCTAGGAA  
GATGATTTATGTTGAAGATAGACTAGCTAAATATATTTCTAGAGTTTGTTATCACTCATTCAAGGTAGTGA  
GAATCTTAAACAGAATTTAGTAGTGAGATATATTCCTGGTGGAGCAAATCAAATAATTTGTAATAATAT  
TTTAACTCATCGTATTTAGATTCGGATAACCATTATTTTTGGCTTGATGGAGATCAAAACACTAATGT  
TAGTGAATCAAATAATTTAATGAATATCTTGAAAAATGGTGTGTTATATCAGATAAAATTCCTGAATC  
AGATAATAAAAAATCTTGATGATATTATAAAATGATAANGGGATGTCCAATTAAATTTAATGTTTCAGG  
TAATAAAGGGCAAAAAAATAATATTGAATTAATTGCGAAACAAAGAAGCTTTATAGATTATTGGGCTAA  
ATAC

**SP100 amino acid (SEQ ID NO:176)**

VNAQSNLILIDEPEISLHPSAIYKFKEFLQECNLKKHQIIITTHSTQLIKDFPREAVKLLVKNGEKV  
DVNIENIDYQDAFFELGDVYHSRKMIYVEDRLAKYILEFVITHSGSENLKQNLVVRYPGGANQIIICNNI  
LNSSYLDSDNHYFWLDGDQNTNVSESNNLMNYLENGVVISDKIPESDNKNLDDIIKLI XGCP I KFNVS G  
NKGQKNNIELIAKQSFIDYWAKY

**SP101 nucleotide (SEQ ID NO:177)**

TTACCGCGTTTCATCAAGATGTCAAACAAGTCATGACCTATCAACCCATGGTGCGAGAAATATTGAGTGA  
ACAAGACACCCCAGCAAACGAAGAGCTTGTGCTTGCTATGATTTATACTGAAACAAAAGGAAAAGAAGG  
CGATGTTATGCAGTCTAGTGAGTCTGCAAGTGGTTCCACCAACACCATCAATGATAATGCCTCTAGCAT  
TCGGCAAGGCATTCAAACCTGACAGGCAATCTCTATCTGGCGCAGAAGAAGGGGGTAGATATCTGGAC  
AGCTGTTCAAGCCTATAATTTTGACCTGCCTATATCGATTTTATCGCCCAAAATGGCAAGGAAAATAC  
CCTGGCTCTAGCCAAACAGTACTCTCGTGAGACTGTTGCCCCCTTGCTTGTAATAGGACTGGAAAGAC  
TTATAGTTATATTACCCCATTTCATTTTTACGGTGCTGAACTCTATGTAAATGGAGGAACTATTA  
TTATTCTAGACAGGTACGACTTAACCTTTACATCATCAAATGTTTCACTCTCTTTTCAACATCTGGC



Table 1

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**SP101 amino acid (SEQ ID NO:178)**

YRVHQDVKQVMYQPMVREILSEQDTPANEELVLAMIYTETKGKEGDMQSSSESASGSTNTINDNASSI  
RQGIQTLTGNLYLAQKKGVDIWTAQYVNFQPAYIDFIAQNGKENTLALAKQYSRETVAPLLGNRTGKT  
YSYIHPISIFHGAELYVNGGNYYSRQVRLNLYIIKCFTLFSTSG

**SP102 nucleotide (SEQ ID NO:179)**

GTGGATGGGCTTTAACTATCTTCGTATTGCGCGTGC GGCTAAAATTGTGGACAATGAGGAGTTTGAAGC  
CTTGATTCTGTACGGGTCAATTGATTGATTGCGCGACCCAGCAGAATTCCACAGAAAACATATCCTTGG  
TGCACGCAATATTCTTTCAAGTCAGTTGAAAAGTAGTCTTGAGCCCTTCGTAAAGATAAACCTGTCCT  
TCTCTACGAAAACCAACGTGCGCAACGAGTTACAAATGCAGCTCTTTACTTGAAAAACAAGGTTTTTC  
TGAGATTTATATCCTTTCTTATGGCTTGGATTCTTGGAAGGGAAAGTGAAGACTAGC

**SP102 amino acid (SEQ ID NO:180)**

WMGFNYLRIRRAAKIVDNEEFELIRTGQLIDLDPAEFHRKHILGARNIPSSQLKTSLAALRKDKPVL  
LYENQRAQRVTNAALYLKKQGFSEIYILSYGLDSWKGVKVKTS

**SP103 nucleotide (SEQ ID NO:181)**

ACTAAACCAGCATCGTTTCGCAGGAAAATAAGGACAATAATCGTGTCTCTTATGTGGATGGCAGCCAGTC  
AAGTCAGAAAAGTGAAAAGTTGACACCAGACCAGGTTAGCCAGAAAGGAATTCAGGCTGAGCAAAT  
TGTAATCAAAAATTACAGATCAGGGCTATGTAACGTCACACGGTGACCACTATCATTACTATAATGGGAA  
AGTTCCCTTATGATGCCCTCTTTAGTGAAGAACTCTTGATGAAGGATCCAAACTATCAACTTAAAGACGC  
TGATATTGTCAATGAAGTCAAGGGTGGTTATATCATCAAGGTCGATGGAAAAATATTATGTCTACCTGAA  
AGATGCAGCTCATGCTGATAATGTTTCAACTAAAGATGAAATCAATCGTCAAAAAACAAGAACATGTCAA  
AGATAATGAGAAGGTTAACTCTAATGTTGCTGTAGCAAGGTCCTCAGGGACGATATACGACAAATGATGG  
TTATGTCTTTAATCCAGCTGATATTATCGAAGATACGGGTAATGCTTATATCGTTTCTCATGGAGGTCA  
CTATCACTACATTTCCCAAGCGATTATCTGCTAGTGAATTAGCAGCAGCTAAAGCACATCTGGCTGG  
AAAAAATATGCAACCGAGTCAGTTAAGCTATTCTTCAACAGCTAGTGACAATAACACGCAATCTGTAGC  
AAAAGGATCAACTAGCAAGCCAGCAATAAATCTGAAAATCTCCAGAGTCTTTTGAAGGAACCTATGTA  
TTCACCTAGCGCCCAACGTTACAGTGAATCAGATGGCCTGGTCTTTGACCCCTGCTAAGATTATCAGTCG  
TACACCAATGGAGTTGCGATTCCGATGGCGACCATTACCACTTTATTCTTTACAGCAAGCTTTCTGCG  
CTTAGAAGAAAAGATTGCCAGAATGGTGCCTATCAGTGGAAGTGGTTCTACAGTTTCTACAAATGCAAA  
ACCTAATGAAGTAGTGTCTAGTCTAGGCAGTCTTTCAAGCAATCCTTCTTCTTTAACGACAAGTAAGGA  
GCTCTCTTCAGCATCTGATGGTTATATTTTTAATCCAAAAGATATCGTTGAAGAAACGGCTACAGCTTA  
TATTGTAAGACATGGTGATCATTTCCATTACATTTCCAAAATCAAATCAAATTGGGCAACCGACTCTTCC  
AAACAATAGTCTAGCAACACCTTCTCCATCTCTTCCAATCAATCCAGGAAGTTCACATGAGAAACATGA  
AGAAGATGGATACGGATTTGATGCTAATCGTATTATCGCTGAAGATGAATCAGGTTTTGTCTATGAGTCA  
CGGAGACCACAATCATTATTTCTTCAAGAA

**SP103 amino acid (SEQ ID NO:182)**

LNQHRSQENKDNRRVSYVDGSQSSQKSENLTDPQVSQKEGIQAEQIVIKITDQGYVTSHGDHYHYNGK  
VPYDALFSEELLMKDPNYQLKDADIVNEVGKGYIIKVDGKYVYVKDAHADNVRTKDEINRQKQEHVK  
DNEKVNNSNVAVARSQGRYTTNDGYVFNPAIDIEDTGNAIVPHGGHYHYIPKSDLSASELAAKAHLAG  
KNMQPSQLSYSSASTADNNTQSVAKGSTSKPANKSENLSLLKELYDSPSAQRYSESDGLVFDPAKIIISR  
TPNGVAIPHGDHYHFIPYSKLSALEEKIARMVPISGTGSTVSTNAKPNEVVSSLGSLSSNPSSLTTSKE  
LSSASDGYIFNPKDIVEETATAYIVRHGDHFHYIPKSNQIGQPTLPNNSLATPSPSLPINPGTSHEKHE  
EDGYGFDANRIIAEDES GFVMSHGDHNHYFFKK

**SP105 nucleotide (SEQ ID NO:183)**

TGACTACCTTGAAATCCCACTTTACAGCTATCTTGGTGGATTCAACACTAAAGTTCTTCCAACCTCCAAT  
GATGAACATCATCAACGGTGGTTCTCACTCTGACGCTCCAATCGCTTTCCAAGAGTTCATGATCTTGCC  
AGTTGGTGCGCCAACATTTAAAGAAGCCCTTCGTACGGTGCTGAAATCTTCCACGCTCTTAAGAAAAT  
CCTTAAATCACGTGGTTTGGAAACTGCCGTAGGTGACGAAGGTGGATTGCTCCTCGTTTTCGAAGGAAC  
TGAAGATGGTGTGTTGAAACTATCCTTGCTGCGATTGAAGCTGCTGGATATGTACCAGGTAAAGACGTATT  
TATCGGATTTGACTGTGCTTCATCAGAATTCTACGATAAAGAACGTAAAGTTTACGACTACACTAAATT  
TGAAGGTGAAGGTGCTGCTGTTTCGTACATCTGCAGAACAAATCGACTACCTTGAAGAATTGGTTAAACAA  
ATACCCAATCATCACTATTGAAGATGGTATGGATGAAAACGACTGGGATGGTTGGAAAGCTCTTACTGA  
ACGTCTTGGTAAGAAAGTACAACTTGTGGTGACGACTTCTTCGTAACAAACACTGACTACCTTGCACG

Table 1

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TGGTATCCAAGAAGGTGCTGCTAACTCAATCCTTATCAAAGTTAACCAAATCGGTACTCTTACTGAAAC  
TTTTGAAGCTATCGAAATGGCTAAAGAAGCTGGTTACACTGCTGTTGTATCACACCGTTCAGGTGAAAC  
TGAAGATTCAACAATCGCTGATATTGCAGTTGCAACTAACGCAGGACAAATCAAGACTGGTTCACCTTC  
ACGTACAGACCGCATCGCTAAATACAACCAATTGCTTCGTATCGAAGACCAACTTGGTGAAAGTAGCTGA  
ATATCGTGGATTGAAATCATTCTACAACCTTAAAAAA

**SP105 amino acid (SEQ ID NO:184)**

DYLEIPLYSYLGGFNTKVLPTPMNLIINGGSHSDAPIAFQEFMILPVGAPTFKEALRYGAEIFHALKKI  
LKSRLGETAVGDEGGFAPRFEGTEDGVETILAAIEAAGYVPGKDVFIGFDCASSEFYDKERKVYDYTKF  
EGEGAAVRTSAEQIDYLEELVNKYPIITIEDGMDENDWDGWKALTERLGKKVQLVGDDFFVTNTDYLAR  
GIQEGAANSILIKVNQIGTLTETFEAIEMAKEAGYTAVVSHRSGETEDSTIADIATNAGQIKTGSLS  
RTDRIAKYNQLLRIEDQLGEVAEYRGLKSFYNLKK

**SP106 nucleotide (SEQ ID NO:185)**

TCGTATCTTTTTTGGAGCAATGTTTCGCGTAGAAGGACATTCCATGGATCCGACCCCTAGCGGATGGCGA  
AATTCTCTTCGTTGTAAAACACCTTCCTATTGACCGTTTTGATATCGTGGTGGCCCATGAGGAAGATGG  
CAATAAGGACATCGTCAAGCGCGTGATTGGAATGCCTGGCGACACCATTTCGTTACGAAAATGATAAACT  
CTACATCAATGACAAAGAAACGGACGAGCCTTATCTAGCAGACTATATCAAACGCTTCAAGGATGACAA  
ACTCCAAAGCACTTACTCAGGCAAGGGCTTTGAAGGAAATAAAGGAACTTTCTTTAGAAGTATCGCTCA  
AAAAGCTCAAGCCTTCACAGTTGATGTCAACTACAACACCAACTTTAGCTTTACTGTTCCAGAAGGAGA  
ATACCTTCTCCTCGGAGATGACCGCTTGGTTTCGAGCGACAGCCGCCACGTAGGTACCTTCAAAGCAAA  
AGATATCACAGGGGAAGCTAAATTCCGCTTATGGCCAATCACCCGTATCGGAACATTT

**SP106 amino acid (SEQ ID NO:186)**

RIFWFSNVRVEGHSMPTLADGEILFVVKHLPIDRFDIVVAHEEDGNKDIVKRVIGMPGDTIRYENDKL  
YINDKETDEPYLADYIKRFKDDKLQSTYSGKGFEGNKGTFFRSIAQKAQAFVTDVNYNTNFSFTVPEGE  
YLLLGDDRLVSSDSRHVGTFKAKDITGEAKFRLWPITRIGTF

**SP107 nucleotide (SEQ ID NO:187)**

GGACTCTCTCAAAGATGTGAAAGCAAATGCTAGCGACAGCAAGCCTGCACAGGACAAGAAGGATGCAAA  
ACAAGGAACGGAAGATAGTAAGGATTAGATAAGATGACTGAAACAAACTCAGTTCCGGCAGGAGTGAT  
TGTGGTCAGTCTACTTGGCCCTCCTAGGCGTGATTGCCCTTCTGGCTGATTCCGCCGTAAGAAAGAGTCAGA  
AATCCAGCAATTAAGCACGGAATTGATCAAGGTTCTAGGACAGCTAGATGCAGAAAAAGCGGATAAAAA  
AGTCCTTGCCAAAGCCCCAAACCTTCTCCAAGAAACCTTGATTTCGTGAAAGAAGAAAATGGCTCAGC  
AGAGACAGAAACTAAACTAGTAGAGGAGCTTAAAGCAATCCTTGACAAACTCAAG

**SP107 amino acid (SEQ ID NO:188)**

DSLKDVKANASDSKPAQDKKDAKQGTEDSKSDKMTETNSVPAGVIVVSLALLGVIAFWLIRRKKESE  
IQQLSTELIKVLGQLDAEKADKKVLAKAQNLLQETLDFVKEENGSAETETKLVEELKAILDKLK

**SP108 nucleotide (SEQ ID NO:189)**

CAAGAAATCCTATCATCTCTTCCAGAAGCAAACAGAGACGAGGGGAATTCAGACTCAGTTGATTGAAGA  
ATCGCTTAGTCAGCAGACTATAATCCAGTCCCTCAATGCTCAAACAGAATTTATCCAAAGATTGCGTGA  
GGCTCATGACAACTACTCAGGCTATTCTCAGTCAGCCATCTTTTATTCTTCAACGGTCAATCCTTCGAC  
TCGCTTTGTAAATGCACTCATTTATGCCCTTTTAGCTGGAGTAGGAGCTTATCGTATCATGATGGGTTT  
AGCCTTGACCGTCGGTCGTTTAGTGACTTTTTTGAAGTATGTTTCAGCAATACACCAAGCCCTTTAACGA  
TATTTCTTCAGTGCTAGCTGAGTTGCAAAGTGCTCTGGCTTGCGTAGAGCGTATCTATGGAGTCTTAGA  
TAGCCCTGAAGTGCTGAAACAGGTAAGGAAGTCTTGACGACCAGTGACCAAGTTAAGGGAGCTATTTT  
CTTTAAACATGTCTCTTTTGGCTACCATCCTGAAAAAATTTGATTAAGGACTTGTCTATCGATATTC  
AGCTGGTAGTAAGGTAGCCATCGTTGGTCCGACAGGTGCTGGAAAATCAACTCTTATCAATCTCCTTAT  
GCGTTTTTATCCCATTAGCTCGGGAGATATCTTGCTGGATGGGCAATCCATTTATGATTATACACGAGT  
ATCATTGAGACAGCAGTTTGGTATGGTGCTTCAAGAAACCTGGCTCACACAAGGGACCATTTCATGATAA  
TATTGCCTTTGGCAATCCTGAAGCCAGTCGAGAGCAAGTAATTGCTGCTGCCAAAGCAGCTAATGCAGA  
CTTTTTTCATCCAACAGTTGCCACAGGGATACGATACCAAGTTGGAAAATGCTGGAGAATCTCTCTCTGT  
CGGCCAAGCTCAGCTCTTGACCATAGCCCAGTCTTTCTGGCTATTCCAAAGATTCTTATCTTAGACGA  
GGCAACTTCTTCCATTGATACACGGACAGAAGTGCTGGTACAGGATGCCTTTGCAAACTCATGAAGGG  
CCGCACAAGTTTCATCATTGCTCACCGTTTGTCAACCATTACAGGATGCGGATTTAATTCTTGTCTTAGT

Table 1

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AGATGGTGATATTGTTGAATATGGTAACCATCAAGAACTCATGGATAGAAAGGGTAAGTATTACCAAAT  
GCAAAAAGCTGCGGCTTTTAGTTCTGA

A

**SP108 amino acid (SEQ ID NO:190)**

KKSYHLFQKQTETRGITQQLIEESLSQQTIIQSFNAQTEFIQRLREAHDNYSQSAIFYSSSTVNPST  
RFVNALIYALLAGVGAYRIMMGSALTVGRLVTFNLNVQQYTKPFNDISSVLAEQLSALACVERIYGVLD  
SPEVAETGKEVLTTSDQVKGAISFKHVSFGYHPEKILIKDLSIDIPAGSKVAIVGPTGAGKSTLINLLM  
RFYPISSGDILLDGQSIYDYTRVSLRQQFGMVLQETWLTQGTIHDNIAFGNPEASREQVIAAAKAANAD  
FFIQQLPQGYDTKLENAGESLSVGQAQLLTIARVFLAIPKILILDEATSSIDTRTEVLVQDAFAKLMKG  
RTSFIIAHLRLSTIQDADLILVLVDGDIVEYGNHQLMDRKGKYYQMKAASFSE

**SP109 nucleotide (SEQ ID NO:191)**

ACGAAATGCAGGGCAGACAGATGCCTCGCAAATTGAAAAGGCGGCAGTTAGCCAAGGAGGAAAAGCAGT  
GAAAAAACAGAAATTAGTAAAGACGCAGACTTGCACGAAATTTATCTAGCTGGAGGTTGTTTCTGGGG  
AGTGGAGGAATATTTCTCACGTGTTCCCGGGGTGACGGATGCCGTTTCAGGCTATGCAAATGGTAGAGG  
AGAAACAACCAAGTACGAATTGATTAACCAACAGGTCATGCAGAAACCGTCCATGTCACCTATGATGC  
CAAGCAAATTTCTCTCAAGGAAATCCTGCTTCACTATTTCCGCATTATCAATCCAACAGCAAAAATAA  
ACAAGGAAATGATGTGGGGACCCAGTACCGTACTGGTGTATTATACACAGATGACAAGGATTTGGAAGT  
GATTAACCAAGTCTTTGATGAGGTGGCTAAGAAATACGATCAACCTCTAGCAGTTGAAAAGGAAAACCTT  
GAAGAATTTTGTGGTGGCTGAGGATTACCATCAAGACTATCTCAAGAAAAATCCAAATGGCTACTGCCA  
TATCAATGTTAATCAGGCGGCCTATCCTGTCATTGATGCCAGCAAATATCCAAAACCAAGTATGAGGA  
ATTGAAAAGACCCTGTACCTGAGGAGTATGCAGTTACCCAGGAAAATCAAACAGAACGAGCTTTCTC  
AAACCGTTACTGGGATAAAATTTGAATCCGGTATCTATGTGGATATAGCAACTGGGGAACCTCTCTTTTC  
ATCAAAAGACAAATTTGAGTCTGGTTGTGGCTGGCCTAGTTTACCCAACCCATCAGTCCAGATGTTGT  
CACCTACAAGGAAGATAAGTCTTACAATATGACGCGTATGGAAGTGCAGGAGCCGAGTAGGAGATTCTCA  
CCTTGGGCATGTCTTTACGGATGGTCCACAGGACAAGGGCGGCTTACGTTACTGTATCAATAGCCTCTC  
TATCCGCTTTATTTCCCAAAGACCAAATGGAAGAAAAAGGCTACGCTTATTTACTAGATTATGTTGAT

**SP109 amino acid (SEQ ID NO:192)**

RNAGQTDASQIEKAAVSQGGKAVKKTEISKDADLHEIYLAGGCFWGVVEEYFSRVPGVTDVSGYANGRG  
ETTKYELINQTHAETVHVITYDAKQISLKEILLHYFRIINPTSKNKQGNVGTQYRTGVYYTDDKDLEV  
INQVFDEVAKKYDQPLAVEKENLKNFVVAEDYHQDYLLKKNPNGYCHINVNQAAYPVIDASKYPKPSDEE  
LKKTLSPEEYAVTQENQTERAFSNRYWDKFESGIYVDIATGEPLFSSKDKFESGCGWPSFTQPISPDVV  
TYKEDKSYNMRMEVRSRVGDSHLGHVFTDGPQDKGLRYCINSLSIRFIPKQMEEEKGYAYLLDYVD

**SP110 nucleotide (SEQ ID NO:193)**

TGTATAGTTTTTACGCTTGTTCTTCTAATTCTGNTAAAAATGAAGAAAATACTTCTAAAGAGCATGCG  
CCTGATAAAATAGTTTTAGATCATGCTTTTCGGTCAAACCTATATTAGATAAAAAACCTGAAAGAGTTGCA  
ACTATTGCTTGGGGAAATCATGATGTAGCATTAGCTTTAGGAATAGTTCCTGTTGGATTTTCAAAGCA  
AATTACGGTGTAAGTGCTGATAAAGGAGTTTTACCATGGACAGAAGAAAAATCAAAGAATAAATGGT  
AAAGCTAACCTATTTGACGATTTGGATGGACTTAACCTTTGAAGCAATATCAAATTCTAAACCAGATGTT  
ATCTTAGCAGGTTATTCTGGTATAACTAAAGAAGATTATGACACTCTATCA

**SP110 amino acid (SEQ ID NO:194)**

CIVFSACSSNSXKNEENTSKEHAPDKIVLDHAFGQTILDKKPERVATIAWGNHDVALALGIVPVGFSKA  
NYGVSADKGVLPWTEEKIKELNGKANLFDLDGLNFEAISNSKPDVILAGYSGITKEDYDTLS

**SP111 nucleotide (SEQ ID NO:195)**

GTGTGTCGAGCATATTCTGAAGCAAACCTATCAAAATATAGAAATTATTTTAGTTGATGACGGTTCTAC  
GGATAATTCTGGGGAAATTTGTGATGCTTTTATGATGCAAGATAATCGTGTGCGAGTATTGCATCAAGA  
AAATAAGGGGGGGCAGCACAAAGCTAAAAATATGGGGATTAGTGTAGCTAAGGGAGAGTACATCACGAT  
TGTTGATTCAGATGATATCGTAAAAGAAAATATGATTGAAACTCTTTATCAGCAAGTCCAAGAAAAGGA  
TGCAGATGTTGTTATAGGGAATTACTATAATTATGACGAAAGTGACGGGAATTTTATTTTATGTAAC  
AGGGCAAGATTTTTGCGTCAAGAATTAGCTATACAAGAAATTATGAACCGTCAAGCAGGAGATTGGAA  
ATTCAATAGCTCGGCCTTTATATTGCCGACATTTAAGTTGATTAAAAAAGAATTATTCAATGAAGTTCA  
CTTTTCAAATGGTTCGCCGCTTTGATGATGAAGCAACTATGCATCGCTTTTATCTTTTAGCCTCTAAAT  
CGTCTTTATAAACGATAATCTCTATCTGTATAGAAGACGTTTCAGGAAGCATCATGAGAACGGAATTTGA

Table 1

TCTTTCTCTGGGCAAGAGATATTGTTGAAGTGTFTTCTAAGAAAAATATCGGATTGTGTCTTGGCTGGTTT  
GGATGTCTCCGTTCTGCGTATTTCGATTTGTCAATCTTTTAAAGATTATAAGCAAACCTTTAGAAATACCA  
TCAATTAACAGATACTGAGGAATATAAAGATATTGTTTCAGATTAAAGTTGTTTTTGTATGCAGAACAA  
AAGAAATGGTAAAAGT

**SP111 amino acid (SEQ ID NO:196)**

CVEHILKQTYQNIIEILVDDGSTDNSGEICDAFMMQDNRRVRLHQENKGGAAQAKNMGISVAKGEYITI  
VSDDDIVKENMIETLYQQVQEKDADVIGNYYNYDESDGNFYFYVTGQDFCVELAIQEIIMNRQAGDWK  
FNSSAFILPTFKLIKKELFNEVHFSNGRRFDDEATMHRFYLLASKIVFINDNLYLYRRRSGSIMRTEFD  
LSWARDIVEVFSKKISDCVLAGLDVSVLRIRFVNLLKDYKQTLFYHQLTDTEYKDICFRLKLFDDAEQ  
RNGKS

**SP0112 nucleotide (SEQ ID NO:197)**

GTGTTTGGATAGCATTTCAGAAATCAGACGTATCAAAATTTTGAGTGTATTATTAATCAATGATGGCTCTCC  
AGATCATTTCATCCAAATATGTGAAGAATTTGTAGAGAAAGATTCTCGTTTCAAATATTTTGAGAAAGC  
AAACGGCGGTCTTTTCATCAGCTCGTAACCTAGGTATTGAATGTTCTGGGGGGGGCGTACATTACTTTTGT  
AGACTCTGATGATTGGTTGGAACATGATGCTTTAGACCGATTATATGGTGCTTTGAAAAAGGAAAACGC  
AGATATTAGTATCGGGCGTTATAATTCTTATGATGAAACACGCTATGTGTATATGACTTATGTTACGGA  
TCCAGATGATTCTCTAGAAGTGATAGAAGGTAAAGCAATTATGGATAGGGAAGGTGTCTGAAGAAGTCAG  
AAATGGGAAGTGGACTGTAGCTGTCTTGAAGTTATTCAAGAGAGAGTTACTACAAGATTTACCATTTC  
TATAGAAAAAATTGCAGAGGATACTTACTGGACATGGAAGGTACTTCTAAGAGCTTCGAGGATAGTCTA  
TTTGAATCGTTGTGTTTACTGGTACCGTGTGGTTTATCTGATACTTTATCGAATACATGGAGTGAAAA  
GCGTATGTATGATGAAATTGGGGCTAGGGAAGAAAAGATAGCTATTTTAGCAAGTTCAGACTATGACTT  
GACCAATCATATTTTGATTTATAAAAAATAGATTACAAAGAGTGATAGCAAAATTAGAAGAACAAAATAT  
GCAGTTCACAGAGATTTACAGAAGAATGATGGAAAAATTGCTTTTACTTCCG

**SP0112 amino acid (SEQ ID NO:198)**

CLDSIQNQTYQNFECLLINDGSPDHSSKICEEFVEKDSRFKYFEKANGGLSSARNLGIIESSGGAYITFV  
DSDDWLEHDAIDRLYGALKKENADISIGRYNSYDETRYVYMTYVTDPPDSLEVI EGKAIMDREGVEEVR  
NGNWTVAVLKLFKRELLQDLFPPIGKIAEDTYWTWKVLLRASRIVYLNRCVYWYRVGLSDTLSNTWSEK  
RMYDEIGAREEKIAILASSDYDLTNHILYKNRLQRVIAKLEEQNMQFTEIYRRMMEKLSLLP

**SP113 nucleotide (SEQ ID NO:199)**

GTGCCTAGATAGTATTATTACTCAAACATATAAAAAATATTGAGATTGTTGTCGTTAATGATGGTTCTAC  
GGATGCTTCAGGTGAAATTTGTAAAGAATTTTCAGAAATGGATCACCGAATTCTCTATATAGAACAAGA  
AAATGCTGGTCTTTCTGCCGCACGAAACACCGGTCTGAATAATATGTCCGGAATTTATGTGACCTTTGT  
GGACTCGGATGATTGGATTGAGCAAGATTATGTAGAACTCTATATAAAAAAATAGTAGAGTATCAGGC  
TGATATTGCAGTTGGTAATTATTATTCTTTCAACGAAAGTGAAGGAATGTTCTACTTTTCATATATTGGG  
AGACTCCTATTATGAGAAAGTATATGATAATGTTTCTATCTTTGAGAACTTGATGAAACTCAAGAAAT  
GAAGAGTTTGTCTTTGATATCTGCTTGGGGTAAACTCTATAAGGCAAGATTGTTTGAGCAGTTGCGCTT  
TGACATAGGTAAATTAGGAGAAGATGGTTACCTCAATCAAAGGTATATTTATTATCAGAAAAGGTAAT  
TTATTTAAATAAAAGTCTTTATGCTTATCGGATTAGAAAAGGTAGTTTATCAAGAGTTTGACAGAAAA  
GTGGATGCACGCTTTAGTTGATGCTATGTCTGAACGTATTACGCTACTAGCTAATATGGGTATCCTCT  
AGAGAAACACTTGGCAGTTTATCGTCAGATGTTGGAAGTCAGTCTCGCCAACGGTCAAGCTAGTGGTTT  
ATCTGACACAGCAACGTATAAAGAGTTTGAAATGAAACAAAGGCTTTTAAATCAGCTATCGAGACAAGA  
GGAAAGTGAAAAGAAAGCCATTGTCCTCGCAGCAAACATATGGCTATGTAGACCAAGTTTAAACGACAAT  
CAAGTCTATTTGTTATCATAATCGTTTCGATTCTGTTTTTATCTGATTTCATAGCGATTTTCCAAATGAATG  
GATTAAGCAATTAAATAAGCGCTTAGAGAAGTTTGACTCAGAAATTATTAATTGTCGGGTAACCTCTGA  
GCAAATTTTCATGTTATAAATCGGATATTAGTTACACAGTCTTTTTACGCTATTTTCATAGCTGATTTCTG  
GCAAGAAGACAAGGCCCTCTACTTGGACTGTGATCTAGTTGTAACGAAAAATCTGGATGACTTGTGTTGC  
TACAGACTTACAAGATTATCCTTTGGCTGCTGTTAGAGATTTTGGGGGCAGAGCTTATTTGGTCAAGA  
AATCTTTAATGCCGTTCTCTTTGGTAAACAATGCTTTTTGGAAAAAGAGAATATGACCCAAAAAATT  
AATTGATGTAACCAATGAATGGCATGATAAGGTGGATCAGGCAGATCAGAGCATCTTGAATATGCTTTT  
TGAACATAAATGTTTGAATTGGACTTTGATTATAATCATATTGTCATTTCATAAACAGTTTGCTGATTA  
TCAATTGCCTGAGGGTCAGGATTATCCTGCTATTATTCACTATCTTTCTCATCGGAAACCGTGGAAAGA  
TTTGGCGGGCCCAACCTATCGTGAAGTTTGGTGGTACTATCATGGGCTTGAATGGACAGAATTGGGACA  
AAACCATCATTTACATCCATTACAAAGATCTCACATCTATCCAATAAAGGAACCTTTCACTTGTCTAAT  
CTATACTGCCTCAGACCATATTGAACAAATTGAGACATTGGTTCAATCCTTGCTGATATTCAGTTTAA

Table 1

GATAGCAGCTAGAGTAATAGTTAGTGATCGATTGGCTCAGATGACAATTTATCCAAACGTGACTATATT  
TAACGGAATTCACATATTTGGTAGATGTGATAATGAATTGGTAGAAACCAGTCAAGTACTTTTAGATAT  
TAATCATGGCGAAAAGACAGAGAAATTTCTCGATCAATTTGCTAATCTTGGCAAGCCTATCTTATCCTT  
TGAAAATACTAAACCTATGAAGTAGGTGAGGAGGCATATGCTGTTGACCAAGTTCAAGCAATGATTGA  
AAAATTGAGAGAAATAAGCAAA

**SP113 amino acid (SEQ ID NO:200)**

CLDSIITQTYKNIEIVVNDGSTDASGEICKEFSEMDHRILYIEQENAGLSAARNTGLNMSGNYVTFV  
DSDDWIEQDYVETLYKKIVEYQADIAVGNYYSFNESEGMFYFHILGDSYIEKVYDNVSIIFENLYETQEM  
KSFALISAWGKLYKARLFEQLRFDIGKLGEDGYLNQKVYLLSEKVIYLNKSLYAYRIRKGSLSRVWTEK  
WMHALVDAMSERITLLANMGYPLEKHLAVYRQMLEVSLANGQASGLSDTATYKEFEMKQRLNQLSRQE  
ESEKKAIVLAANYGYVDQVLTITKSICYHNRSIRFYLIHSDFPNEWIKQLNKRLEKFDSEIINCRVTSE  
QISCYKSDISYTVFLRYFIADFVQEDKALYLDCLDVVTKNLDDLFDLQDYPLAAVRDFGGRAYFGQE  
IFNAGVLLVNNAFWKENMTQKLIDVTNEWHDKVDQADQSILNMLFEHKWLELDFDYNHIVHKQFADY  
QLPEGQDYPPIIHLYSHRKPWKDLAAQTYREVWYYHGLEWTELQNHHLHPLQRSHIYPIKEPFTCLI  
YTASDHIEQIETLVQSLPDIQFKIAARVIVSDRLAQMTIYPNVTIFNGIHYLVDVDNELVETSQVLLDI  
NHGEKTEEILDQFANLGKPILSFENTKTYEVGQEAAYVDQVQAMIEKLREISK

**SP114 nucleotide (SEQ ID NO:201)**

CATTGAGAAGCAGACCTATCAAAATCTGGAAATATTCTTGTGATGATGGTGCAACAGATGAAAGTGG  
TCGCTTGTGTGATTCAATCGCTGAACAAGATGACAGGGTGTGAGTGCCTTCATAAAAAGAACGAAGGATT  
GTCGCAAGCACGAAATGATGGGATGAAGCAGGCTCACGGGGATTATCTGATTTTTATTGACTCAGATGA  
TTATATCCATCCAGAAATGATTCAGAGCTTATATGAGCAATTAGTTCAGAAGATGCCGATGTTTCGAG  
CTGTGGTGTGATGAATGTCTATGCTAATGATGAAAGCCACAGTCAGCCAATCAGGATGACTATTTTGT  
CTGTGATTCTCAACACATTTCTAAAGGAATACCTCATAGGTGAAAAAATACCTGGGACGATTGCAATAA  
GCTAATCAAGAGCAGATGTCAGTGCCTATCCTTTCTTAAGGGGTTGATTTACGAAGATGCCTATTA  
CCATTTTGTATTTAATCAAGTTGGCCAAGAAGTATGTGGTTAATACTAAACCCTATTATTACTATTTTCCA  
TAGAGGGGATAGTATTACGACCAAACCCCTATGCAGAGAAGGATTTAGCCTATATTGATATCTACCAAAA  
GTTTTATAATGAAGTTGTGAAAACTATCCTGACTTGAAAGAGGTGCGCTTTTTTCAGATTGGCCTATGC  
CCACTTCTTTTATTCTGGATAAGATGTTGCTAGATGATCAGTATAAACAGTTTGAAGCCTATTCCTCAGAT  
TCATCGTTTTTTTAAAGGCCATGCCTTTGCTATTCTAGGAATCCAATTTTCCGTAAGGGGAGAAGAAT  
TAGTGCTTTGGCCCTATTTCATAAATATTTCTTATATCGATTCTTATTACTGAAAAATATTGAAAAATC  
TAAAAAATTACAT

**SP114 amino acid (SEQ ID NO:202)**

IQKQTYQNLEIILVDDGATDESGRLCDSIAEQDDRVSVLHKKNEGLSQARNDGMKQAHGDYLIIFIDSD  
YIHPEMIQSLYEQLVQEDADVSSCGVMNVYANDESPQSANQDDYFVCDQSQTFLKEYLIGEKIPGTICNK  
LIKRIATALSFPKGLIYEDAYYHFDLIKLAKKYVNTKPYYYYFHRGDSITTKPYAEKDLAYIDIYQK  
FYNEVVKNYPDLKEVAFFRLAYAHFFILDKMLDDQYKQFEAYSQIHRFLKGHAF AISRNPIFRKGRRI  
SALALFINISLYRFLLLKNIEKSKKLH

**SP115 nucleotide (SEQ ID NO:203)**

TAAGGCTGATAATCGTGTTCAAATGAGAACGACGATTAATAATGAATCGCCATTGTTGCTTTCTCCGTT  
GTATGGCAATGATAATGGTAACGGATTATGGTGGGGGAACACATTGAAGGGAGCATGGGAAGCTATTCC  
TGAAGATGTAAAGCCATATGCAGCGATTGAACTTCATCCTGCAAAAGTCTGTAAACCAACAAGTTGTAT  
TCCACGAGATACGAAAGAATTGAGAGAATGGTATGTCAAGATGTTGGAGGAAGCTCAAAGTCTAAACAT  
TCCAGTTTTCTTGTTATTATGTCGGCTGGAGAGCGTAATACAGTTCCTCCAGAGTGGTTAGATGAACA  
ATTCCAAAAGTATAGTGTGTTAAAGGTGTTTTAAATATTGAGAATTATTGGATTTACAATAACCAGTT  
AGCTCCGCATAGTGCTAAATATTTGGAAGTTTGTGCCAAATATGGAGCGCATTTTATCTGGCATGATCA  
TGAAAAATGGTTCTGGGAACTATTATGAATGATCCGACATTCTTTGAAGCGAGTCAAAAATATCATAA  
AAATTTGGTGTGGCAACTAAAAATACGCCAATAAGAGATGATGCGGGTACAGATTCTATCGTTAGTGG  
ATTTTGGTTGAGTGGCTTATGTGATAACTGGGCTCATCAACAGATACATGGAAATGGTGGGAAAAACA  
TTATACAAACACATTTGAACTGGAAGAGCTAGGGATATGAGATCCTATGCATCGGAACCAGAATCAAT  
GATTGCTATGGAAATGATGAATGTATATACTGGGGGAGGCACAGTTTATAATTTCGAATGTGCCCGGTA  
TACATTTATGACAAATGATGTACCAACTCCAGCATTTACTAAAGGTATTATTCTTTCTTTAGACATGC  
TATACAAAATCCAGCTCCAAGTAAGGAAGAAGTTGTAAATAGAACAAAAGCTGTATTTTGGAAATGGAGA  
AGGTAGGATTAGTTCATTAAACGGATTTTATCAAGGACTTTATTTCGAATGATGAAACAATGCCTTTATA  
TAATAATGGGAGATATCATATTCTTCTGTAATACATGAGAAAATTGATAAGGAAAAGATTTTCATCTAT

Table 1

ATTCCCTAATGCAAAAATTTTACTAAAAATAGTGAGGAATTGTCTAGTAAAGTCAACTATTTAAACTC  
GCTTTATCCAAAACCTTTATGAAGGAGATGGGTATGCTCAGCGTGTAGGTAATTCCTGGTATATTTATAA  
TAGTAATGCTAATATCAATAAAAAATCAGCAAGTAATGTTGCCATGTATATACTAATAATACAAAGTCGTT  
ATCGTTAGATTTGACGCCACATACTTACGCTGTTGTTAAAGAAAATCCAAATAATTTACATATTTTATT  
GAATAATTACAGGACAGATAAGACAGCTATGTGGGCATTATCAGGAAAATTTGATGCATCAAAAAGTTG  
GAAGAAAGAAGAATTAGAGTTAGCGAACTGGATAAGCAAAAATTTATCCATCAATCCTGTAGATAATGA  
CTTTAGGACAACAACACTTACATTAAGGGGCATCTGGTCATAAACCTCAGATAAATATAAGTGGCGA  
TAAAAATCATTATACTTATACAGAAAATTTGGGATGAGAATACCCATGTTTATACCATTACGGTTAATCA  
TAATGGAATGGTAGAGATGTCTATAAATACGTAGGGGACAGGTCCAGTCTCTTTCCCAACACCAGATAA  
ATTTAATGATGGTAATTTGAATATAGCATATGCAAAACCAACACAAAGTTCTGTAGATTACAATGG  
AGACCCTAATAGAGCTGTGGATGGTAACAGAAATGGTAATTTTAACTCTGGTTCGGTAACACACACTAG  
GGCAGATAATCCCTCTTGGTGGGAAGTCGATTTGAAAAAATGGATAAAGTTGGGCTTGTTAAAATTTA  
TAATCGCACAGATGCTGAGACTCAACGTCTATCTAATTTT

**SP115 amino acid (SEQ ID NO:204)**

KADNRVQMRTTINNESPLLLSPLYGNDNGNGLWWGNTLKGAWAIPEDVKPYAAIELHPAKVCKPTSCI  
PRDTKELREWYVKMLEEAQSLNIPVFLVIMSAGERNTVPPEWLDEQFQKYSVLKGVLNINIENYWIYNNQL  
APHSAKYLEVCAKYGAHFIWHDHEKWFWEITMNDPTFFEASQKYHKNLVLATKNTPIRD DAGTDSIVSG  
FWLSGLCDNWGSSDTWKWWEKHNTFTETGRARDMRSYASEPESMIAMEMMNVTYGGGTVYNFECAAY  
TFMTNDVPTPAFTKGIIPFFRHAIQNPAPSKEEVNRTKAVFWNGEGRISSLNGFYQGLYSNDETPLY  
NNGRYHILPVIHEKIDKEKISSIFPNAKILTKNSEELSSKVNLYNSLYPKLYEGDGYAQRVGNWYIYN  
SNANINKNQVMLPMYTNNTKSLSLDLTPHTYAVVKENPNLHILLNMYRTDKTAMWALSGNFDASKSW  
KKEELELANWISKYNSINPVDNDFRTTTLTKGHTGHKPQINISGDKNHYTYTENWDENTHVTITVNH  
NGMVEMSIINTEGTGPVVSFTPDKFNDGNLNIAYAKPTTQSSVDYNGDPNRAVDGNRNGNFNSGSVTHTR  
ADNPSWWEVDLKKMDKVGLVKIYNRTDAETQRLSNF

**SP117 nucleotide (SEQ ID NO:205)**

CTGTGGCAATCAGTCAGCTGCTTCCAAACAGTCAGCTTCAGGAACGATTGAGGTGATTTACAGAGAAAA  
TGGCTCTGGGACACGGGGTGCCTTCACAGAAATCAGAGGATTCTCAAAAAAGACGGTGATAAAAAAAT  
TGACAACACTGCCAAACAGCTGTGATTCAAAATAGTACAGAAGGTGTTCTCTCAGCAGTTCAAGGGAA  
TGCTAATGCTATCGGCTACATCTCCTTGGGATCTTTAACGAAATCTGTCAAGGCTTTAGAGATTGATGG  
TGTCAGGCTAGTCGAGACACAGTTTTAGATGGTGAATACCCCTCTTCAACGTCCCTTCAACATTGTTTG  
GTCTTCTAATCTTTCCAAGCTAGGTCAAGATTTTATCAGCTTTATCCACTCCAAACAAGGTCAACAAGT  
GGTCACAGATAATAAATTTATTGAAGCTAAACCCGAAACCACGAATATACAAGCCAACACTTATCAGG  
CAAGTTGTCTGTTGTAGGTTCCACTTCAGTATCTTCTTAAATGGAAAAATTAGCAGAAGCTTATAAAAA  
AGAAAAATCCAGAAGTTACGATTGATATTACCTCTAATGGGTCTTCAGCAGGTATTACCGCTGTTAAGGA  
GAAACCGCTGATATTGGTATGGTTTCTAGGGAAATTAACCTCTGAAGAAGGTAAGAGTCTCACCCATGA  
TGCTATTGCTTTAGACGGTATTGCTGTTGTGGTCAATAATGACAATAAGGCAAGCCAAGTCAGTATGGC  
TGAACCTGCAGACGTTTTTAGTGGCAAATTAACCACCTGGGACAAGATTAAA

**SP117 amino acid (SEQ ID NO:206)**

CGNQSAASKQSASGTIEVISRENGSGTRGAFTEITGILKKDGDKKIDNTAKTAVIQNSTEGVLSAVQGN  
ANAIGYISLGLTKSVKALEIDGVKASRDVLDGEYPLQRPFNIVWSSNLSKLGQDFISFIHKSQGGQV  
VTDNKFIEAKTETTEYTSQHLVGKLSVVGSTSVSSLMEKLAELAYKKENPEVTIDITSNGSSAGITAVKE  
KTADIGMVSRELTPPEGKSLTHDAIALDGIADVNNNDNKASQVSMALADVFSGLKLTWDKIK

**SP118 nucleotide (SEQ ID NO:207)**

TTGTCAACAACAACATGCTACTTCTGAGGGGACGAATCAAAGGCAAAGCAGTTCAGCGAAAGTTCCATG  
GAAAGCTTCATACACCAACCTAAACAACCAGGTAAGTACAGAAGAGGTCAAATCTCTCTTATCAGCTCA  
CTTGGATCCAAATAGTGTGATGCATTTTTTAATCTCGTTAATGACTATAATACCATTGTCGGCTCAAC  
TGGCTTATCAGGAGATTTCACTTCCTTTACTCACACCGAATACGATGTTGAGAAAATCAGTCATCTCTG  
GAATCAAAAAGAAGGGCGATTTTGTGGGACCAACTGCCGTATCAATAGTTATTGTCTTTTGAAAAATTC  
AGTCACCATTCCAAAGCTTGAAAAGAAATGACCAGTTGCTTTTCTAGATAATGATGCGATTGATAAAGG  
AAAGGTCTTTGATTACACAAGATAAGGAAGAGTTTGATATTCTATTTTCGAGAGTTCCAACCTGAGTCAAC  
TACAGATGTCAAGGTTACGCTGAAAAGATGGAAGCATTCTTCTCACAATTTCAATTCAATGAAAAAGC  
TCGAATGCTGTCTGTAGTCTTGACGACAATTTGGATGGCGAGTATCTGTTGTAGGCCACGTTGGGGT  
CTTAGTACCTGCTGATGACGGTTTCTTATTTGTAGAGAAATTGACTTTCGAAGAGCCCTACCAAGCGAT

Table 1

TAAATTTGCTAGTAAGGAAGATTGCTACAAGTATTTGGGCACCAAGTATGCGGATTATACAGGCGAGGG  
ACTGGCTAAGCCTTTTATCATGGATAATGATAAGTGGGTTAAACTT

**SP118 amino acid (SEQ ID NO:208)**

CQQQHATSEGTNQRQSSSAKVPWKASYTNLNNQVSTEEVKSLLSAHLDPNSVDAFFNLVNDYNTIVGST  
GLSGDFTSFTHTEYDVEKISHLWNQKKGDFVGTNCRINSYCLLKNSVTIPKLEKNDQLLFLDNDIDAIDKG  
KVFDSQDKKEFDILFSRVPTESTTDVKVHAEKMEAFFSQFQFNEKARMLSVVLHDNLDGEYLFVGHVGV  
LVPADDGFLFVEKLTFEOPYQAIKFASKEDCYKYLGTQYADYTGEGLAKPFI MDNDKWVKL

**SP119 nucleotide (SEQ ID NO:209)**

TTGTTTCAGGCAAGTCCGTGACTAGTGAACACCAAGATGAAATGAAGACGGAGCAGACAGCTAG  
TAAAACAAGCGCAGCTAAAGGGAAAGAGTGGCTGATTTTGAATTGATGGGAGTAGATGGCAAGACCTA  
CCGTTTATCTGATTACAAGGGCAAGAAAGTCTATCTCAAATTCTGGGCTTCTTGGTGTTCATCTGTCT  
GGCTAGTCTTCCAGATACGGATGAGATTGCTAAAGAAGCTGGTGATGACTATGTGGTCTTGACAGTAGT  
GTCACCAGGACATAAGGGAGAGCAATCTGAAGCGGACTTTAAGAATTGGTATAAGGGATTGGATTATAA  
AAATCTCCCAGTCTAGTTGACCCATCAGGCAAACTTTTGAAACTTATGGTGTCCGTTCTTACCCAAC  
CCAAGCCTTTATAGACAAAAGGCAAGCTGGTCAAAACACATCCAGGATTCATGGAAAAAGATGCAAT  
TTTGCAAACTTTGAAGGAATTAGCC

**SP119 amino acid (SEQ ID NO:210)**

CSGKSVTSEHQTKDEMKTETASKTSAKKGKEVADFELMGVDGKTYRLSDYKGGKVVYLFKFWASWCSICL  
ASLPDTDEIAKEAGDDYVVLTVVSPGHKGEQSEADFKNWKGLDYKNLPVLVDPGKLLLETYGVRSYPT  
QAFIDKEGKLVKTHPGFMEKDAILQTLKELA

**SP120 nucleotide (SEQ ID NO:211)**

CTCGCAAATTGAAAAGGCGGCAGTTAGCCAAGGAGGAAAAGCAGTGAAAAAACAGAAATTAGTAAAGA  
CGCAGACTTGACAGAAATTTATCTAGCTGGAGGTTGTTTCTGGGGAGTGGAGGAATATTTCTCACGTGT  
TCCCGGGGTGACGGATGCCGTTTCAGGCTATGCAAATGGTAGAGGAGAAACAACCAAGTACGAATTGAT  
TAACCAAAACAGGTCTACGAGAAACCGTCCATGTACCTATGATGCCAAGCAAATTTCTCTCAAGGAAAT  
CCTGCTTCACTATTTCCGCAATTATCCAACAGCAAAAATAACAAGGAAATGATGTGGGGACCCA  
GTACCGTACTGGTGTGTTTATTACACAGATGACAAAGGATTTGGAAGTGATTAACCAAGTCTTTGATGAGGT  
GGCTAAGAAATACGATCAACCTCTAGCAGTTGAAAAGGAAAACTTGAAGAATTTGTGGTGGCTGAGGA  
TTACCATCAAGACTATCTCAAGAAAAATCCAATGGCTACTGCCATATCAATGTTAATCAGGCGGCCTA  
TCCTGTCAATTGATGCCAGCAAATATCCAAAACCAAGTATGAGGAATTGAAAAAGACCCTGTCACTGA  
GGAGTATGCAGTTACCCAGGAAAAATCAAACAGAACGAGCTTTCTCAAACCGTTACTGGGATAAATTTGA  
ATCCGGTATCTATGTGGATATAGCAACTGGGGAACCTCTCTTTTCATCAAAAAGACAAATTTGAGTCTGG  
TTGTGGCTGGCCTAGTTTTTACCCAACCCATCAGTCCAGATGTTGTCACCTACAAGGAAGATAAGTCCTA  
CAATATGACGCGTATGGAAGTGCGGAGCCGAGTAGGAGATTCTCACCTTGGGCATGTCTTTACGGATGG  
TCCACAGGACAAGGGCGGCTTACGTTACTGTATCAATAGCCTCTCTATCCGCTTTATTCCCAAAGACCA  
AATGGAAGAAAAAGGTACGCTTATTTAC

**SP120 amino acid (SEQ ID NO:212)**

SQIEKAAVSQGGKAVKKTEISKDADLHEIYLAGGCFWGVVEYFSRVPGVTDVAVSGYANGRGETTKYELI  
NQTGHAETVHVITYDAKQISLKEILLHYFRIINPTSKNKQGNVGTQYRTGVYYTDDKDLEVINQVFDEV  
AKKYDQPLAVEKENLKNFVVAEDYHQDYLLKNPNNGYCHINVNQAAYPVIDASKYPKPSDEELKKTLSPE  
EYAVTQENQTERAFSNRYWDKFESGIYVDIATGEPLFSSKDKFESGCGWPSFTQPI SPDVVITYKEDKSY  
NMTRMEVRSRVGDSHLGHVFTDGPQDKGLRYCINSLSIRFIPKQDMEEKGTLIY

**SP121 nucleotide (SEQ ID NO:213)**

TTGTCAGTCAGGTTCTAATGGTTCTCAGTCTGCTGTGGATGCTATCAAAACAAAAGGGAAATTAGTTGT  
GGCAACCAGTCCTGACTATGCACCTTTGAATTTCAATCATTTGGTTGATGGAAAGAACCAGGTAGTCGG  
TGCAGACATCGACATGGCTCAGGCTATCGCTGATGAACCTGGGGTTAAGTTGGAAATCTCAAGCATGAG  
TTTGTACAATGTTTTGACCAGTCTTCAAACCTGGTAAGGCTGACCTAGCAGTTGCAGGAATTAGTGCTAC  
TGACGAGAGAAAAAGTCTTTGATTTTTTCAATCCCATACTATGAAAACAAGATTAGTTTCTTGGTTTCG  
TAAGGCTGATGTGGAAAAATACAAGGATTTAACTAGCCTAGAAAGTGCTAATATTGCAGCCCCAAAAGG  
GACTGTTCCAGAATCAATGGTCAAGGAACAATTGCCAAAAGTTCAATTAACCTCCCTAACTAATATGGG  
TGAAGCAGTCAATGAATTGCAGGCTGGAAAAATAGATGCTGTTTCATATGGATGAGCCTGTTGCACCTAG

Table 1

92

TTATGCTGCTAAAAACGCTGGCTTAGCTGTGCGAACTGTCAGCTTGAAGATGAAGGACGGCGACGCCAA  
TGCC

**SP121 amino acid (SEQ ID NO:214)**

CQSGSNGSQSAVDAIKQKGLVVATSPDYAPFEFQSLVDGKNQVVGADIDMAQAIADDELGVKLEISSMS  
FDNVLTSLQTKADLAVAGISATDERKEVDFDSIPYYENKISFLVRKADVEKYKDLTSLESANIAAQKG  
TVPESMVKEQLPKVQLTSLTNMGEAVNELQAGKIDAVHMDPEVALSYAKNAGLAVATVSLKMKDKGDAN  
A

**SP122 nucleotide (SEQ ID NO:215)**

GGAACTTCACAGGATTTTAAAGAGAAGAAAACAGCAGTCATTAAGGAAAAAGAAGTTGTTAGTAAAAA  
TCCTGTGATAGACAATAACACTAGCAATGAAGAAGCAAAAATCAAAGAAGAAAATTCGAATAAATCCCA  
AGGAGATTATACGGACTCATTTGTGAATAAAAACACAGAAAATCCCAAAAAGAAGATAAAGTTGTCTA  
TATTGCTGAATTTAAAGATAAAGAATCTGGAGAAAAAGCAATCAAGGAAGTATCCAGTCTTAAGAATAC  
AAAAGTTTATATACTTATGATAGAATTTTAAACGGTAGTGCCATAGAAAACAAGTCCAGATAACTTGGGA  
CAAAATTAACAATAAGAGGTATTTTCATCGGTTGAAAGGGCACAAAAGTCCAACCCATGATGAATCA  
TGCCAGAAAAGGAAATTGGAGTTGAGGAAGCTATTGATTACCTAAAGTCTATCAATGCTCCGTTTGGGAA  
AAATTTTGATGGTAGAGGTATGGTCATTTCAAATATCGATACTGGAACAGATTATAGACATAAGGCTAT  
GAGAATCGATGATGATGCCAAAGCCTCAATGAGATTTAAAAAAGAAGACTTAAAGGCACTGATAAAAA  
TTATTGGTTGAGTGATAAAATCCCTCATGCGTTCAATTATTATAATGGTGGCAAAATCACTGTAGAAAA  
ATATGATGATGGAAGGGATTATTTTGACCCACATGGGATGCATATTGCAGGGATTCTTGCTGGAAATGA  
TACTGAACAAGACATCAAAAACCTTTAACGGCATAGATGGAATTGCACCTAATGCACAAATTTTCTCTTA  
CAAAATGTATTCTGACGCAGGATCTGGGTTTGGCGGTGATGAAACAATGTTTCATGCTATTGAAGATTC  
TATCAAAACACAACGTTGATGTTGTTTCGGTATCATCTGGTTTACAGGAACAGGTCTTGTAGGTGAGAA  
ATATTGGCAAGCTATTCCGGCATTAAGAAAAGCAGGCATTCCAATGGTTGTCGCTACGGGTAAGTATGC  
GACTTCTGCTTCAAGTTCTTCATGGGATTTAGTAGCAAATAATCATCTGAAAATGACCGACACTGGAAA  
TGTAACACGAACTGCAGCACATGAAGATGCGATAGCGGTGCTTCTGCTAAAAATCAAACAGTTGAGTT  
TGATAAAGTTAACATAGGTGGAGAAAGTTTAAATACAGAAATATAGGGGCTTTTTCGATAAGAGTAA  
AATCACAACAAATGAAGATGGAACAAAAGCTCCTAGTAAATTAATTTGTATATATAGGCAAGGGGCA  
AGACCAAGATTGTATAGGTTTGGATCTTAGGGGCAAAATGCAGTAATGGATAGAATTTATACAAAGGA  
TTTAAAAAATGCTTTTAAAAAAGCTATGGATAAGAGGTGCACGCGCCATTATGGTTGTAAATACTGTAAA  
TTACTACAATAGAGATAATTGGACAGAGCTTCCAGCTATGGGATATGAAGCGGATGAAGGTACTAAAAG  
TCAAGTGTTTCAATTTTCAGGAGATGATGGTGTAAGCTATGGAAACATGATTAATCCTGATAAAAAAAC  
TGAAGTCAAAAAGAAATAATAAAGAAGATTTTAAAGATAAATTTGGAGCAATACTATCCAATTGATATGGA  
AAGTTTAAATTCACAACAAACCGAATGTAGGTGACGAAAAAGAGATTGACTTTAAGTTTGCACCTGACAC  
AGACAAAGAACTCTATAAAGAAGATATCATCGTTCCAGCAGGATCTACATCTTGGGGGCCAAGAATAGA  
TTTACTTTTAAACCCGATGTTTCAGCACCTGGTAAAAATATTAAATCCACGCTTAATGTTATTAATGG  
CAAATCAACTTATGGCTATATGTCAGGAAC TAGTATGGCGACTCCAATCGTGGCAGCTTCTACTGTTTT  
GATTAGACCGGAAATTAAGGAAATGCTTGAAAGACCTGTATTGAAAAATCTTAAGGGAGATGACAAAAT  
AGATCTTACAAGTCTTACAAAATTTGCCCTACAAAATACTGCGCGACCTATGATGGATGCAACTTCTTG  
GAAAGAAAAAAGTCAATACTTTGCATCACCTAGACAACAGGGAGCAGGCCTAATTAATGTGGCCAATGC  
TTTGAGAAAATGAAGTTGTAGCAACTTTCAAAAACACTGATTCTAAAGGTTTGGTAAACTCATATGGTTC  
CATTTCTCTTTAAAGAAATAAAAGGTGATAAAAAATACTTTACAATCAAGCTTCACAATACATCAAACAG  
ACCTTTGACTTTTAAAGTTTCAGCATCAGCGATAACTACAGATTCTCTAACTGACAGATTAAACTTGA  
TGAAACATATAAAGATGAAAAATCTCCAGATGGTAAGCAAAATTTGTTCCAGAAATTCACCCAGAAAAAGT  
CAAAGGAGCAAATATCACATTTGAGCATGATACTTTCACTATAGGCGCAAATTTCTAGCTTTGATTGAA  
TGCGGTTATAAATGTTGGAGAGGCCAAAAACAAAAATAAATTTGTAGAATCATTTATTCATTTTGAGT  
AGTGGAAGCGATGGAAGCTCTAAACTCCAGCGGGAAGAAAAATAAATTTCCAACCTTCTTTGTCGATGCC  
TCTAATGGGATTTGCTGGGAATTGGAACACGGAACCAATCTTGATAAATGGGCTTGGGAAGAAGGGTC  
AAGATCAAAAACACTGGGAGGTTATGATGATGATGGTAAACCGAAAAATTCAGGAACCTTAAATAAGGG  
AATTTGGTGGAGAACATGGTATAGATAAATTTAATCCAGCAGGAGTTATACAAAATAGAAAAGATAAAAA  
TACAACATCCCTGGATCAAAATCCAGAATTATTTGCTTTCAATAACGAAGGGATCAACGCTCCATCATC  
AAGTGGTTCTAAGATTGCTAACATTTATCCTTTAGATTCAAAATGGAAATCCTCAAGATGCTCAACTTGA  
AAGAGGATTAACACCTTCTCCACTTGTATTAAGAAGTGCAGAAGAAGGATTGATT

**SP122 amino acid (SEQ ID NO:216)**

ETSQDFKEKKTAVIKEKEVSKNPVIDNNTSNEEAKIKEENSNSQGDYTDSEFVNKNTENPKKEDKVY  
IAEFKDKESGEKAIKELSSLKNTKVLYTYDRIFNGSAIETTPDNLDKIKQIEGISSVERAQKVQPMNHN

006634-0300  
102270-123920



Table 1

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ARKEIGVEEAIDYLSINAPFGKNFDGRGMVISNIDTGTDRHKAMRIDDDAKASMRFKKEDLKGTDKN  
 YWLSDKIPHAFNYNGGKITVEKYDDGRDYDFDPHGMHIAGILAGNDTEQDIKNFNGIDGIAPNAQIFSY  
 KMYSDAGSGFAGDETMFHAIEDSIKHNVDVSVSSGFTGTGLVGEKYWQAIRALRKAGIPMVVATGNYA  
 TSASSSSWDLVANNHLKMTDTGNVTRTAAHEDAIVASAKNQTFEFDKVNIGGESFKYRNIGAFFDKSK  
 ITTNEDEGTKAPSKLKFVYIGKGQDQDLIGLDLRGKIAVMDRIYTKDLKNAFKKAMDKGARAIMVVNTVN  
 YYNRDNWTELPAMGYEADGKTSQVFSISGDDGVKLWNMINPDKKTEVKRNKEDFKDKLEQYYPIDME  
 SFNSNKPNGVDEKEIDFKFAPDTEKELYKEDIIVPAGSTSWGPRIDLKPDVSAPGKNIKSTLNVING  
 KSTYGYMSGTSMATPIVAASTVLIRPKLKEMLERPVLKNLKGDDKIDLTSLTKIALQNTARPMMDATSW  
 KEKSQYFASPRQGGAGLINVANALRNEVVATFKNTDSKGLVNSYGSISLKEIKGDKKYFTIKLHNTSNR  
 PLTFKVSASAITTDSLTDRLKLDETYKDEKSPDGKQIVPEIHPEKVKGANITFEHDTFTIGANSSFDLN  
 AVINVGEAKNKNKFVESFIHFESVEAMEALNSSGKKINFQPSLSMPLMGFAGNWNHEPILDKWAVEEGS  
 RSKTLGGYDDDGKPKIPGTLNKGIGGEHGIDKFNPAQVQNRKDKNTTSLDQNPFLFAFNNEGINAPSS  
 SGSKIANIYPLDSNGNPQDAQLERGLTPSPLVLRSAEEGLI

**SP123 nucleotide (SEQ ID NO:217)**

TGTGGTCGAAGTTGAGACTCCTCAATCAATAACAAATCAGGAGCAAGCTAGGACAGAAAACCAAGTAGT  
 AGAGACAGAGGAAGCTCCAAAAGAAGAAGCACCTAAAACAGAAGAAAGTCCAAAAGGAAGAACCAAAATC  
 GGAGGTAACCTACTGACGACACCCCTTCTTAAAGTAGAAGAGGGGAAAGAAGATTGAGCAGAACCCAGC  
 TCCAGTTGAAGAAGTAGGTGGAGAAGTTGAGTCAAAACCAGAGGAAAAAGTAGCAGTTAAGCCAGAAAG  
 TCAACCATCAGACAAACCAGCTGAGGAATCAAAAGTTGAACAAGCAGGTGAACAGTCGCGCCAAGAGA  
 AGACGAAAAAGGCACCAAGTCGAGCCAGAAAAGCAACCAGAAGCTCCTGAAGAAGAGAAGGCTGTAGAGGA  
 AACACCGAAAAACAAGAAGAGTCAACTCCAGATACCAAGGCTGAAGAACTGTAGAACCAAAAGAGGAGAC  
 TGTTAATCAATCTATTGAACAACCAAAAGTTGAAACGCCTGCTGTAGAAAAACAAAACAGAACCAACAGA  
 GGAACCAAAAGTTGAACAAGCAGGTGAACCAAGTCGCGCCAAGAGAAGACGAACAGGCACCAACGGCACC  
 AGTTGAGCCAGAAAAGCAACCAGAAGTTCCTGAAGAAGAGAAGGCTGTAGAGGAAAACACCGAAACAGAA  
 AGATAAAATAAAGGGTATTGGTACTAAAGAACCAGTTGATAAAAGTGAGTTAAATAATCAAATTGATAA  
 AGCTAGTTTCACTTCTCCTACTGATTATTCTACAGCAAGTTACAATGCTCTTGGACCTGTTTTAGAAAC  
 TGCAAAAGGTGTCTATGCTTTCAGAGCCTGTAAACAGCCTGAGGTAAATAGCGAGACAAATAAACTTAA  
 AACCGCTATTGACGCTCTAAACGTTGATAAAAGTGAATTAACAATACGATTGCAGATGCAAAAACAAA  
 GGTAAGAAGACATTACAGTGATGAAGTTGGCAAAACCTCCAACTGAAGTTACAAAGGCTGAAAAGGT  
 TGCAGCTAATACAGATGCTAAACAAAGTGAAGTTAACGAAGCTGTTGAAAAATTAAGTGAACATTTGA  
 AAAATTGGTTGAATTATCTGAAAAGCCAATATTAACATTGACTAGTACCGATAAGAAAAATATTGGAACG  
 TGAAGCTGTTGCTAAGTATACTCTAGAAAATCAAAACAAAACAAAAATCAAATCAATCACAGCTGAATT  
 GAAAAAAGGAGAAGAAGTTATTAATACTGTAGTCTTACAGATGACAAGGTAACAACAGAACTATAAG  
 CGCTGCATTTAAGAACCTAGAGTACTACAAAGAATACACCCTATCTACAACATGATTTACGACAGAGG  
 TAACGGTGAAGAACTGAACTCTAGAAAATCAAAATATTCAATTAGATCTTAAAAAAGTTGAGCTTAA  
 AAATATTAAACGTACAGATTTAATCAAATACGAAAATGGAAGAACTAATGAATCACTGATAACAAC  
 TATTCCTGATGATAAGAGCAATTATTATTTAAAAATAACTTCAAATAATCAGAAAACCTACATTACTAGC  
 TGTTAAAAATATAGAAGAACTACGGTTAACGGAACACCTGTATATAAAGTTACAGCAATCGCAGACAA  
 TTTAGTCTCTAGAAGTGTGATAATAAATTTGAAGAAGAA

**SP123 amino acid (SEQ ID NO:218)**

VVEVETPQSITNQEQARTENQVVETEEAPKEEAPKTEESPKEEPKSEVKPTDDTLPKVEEGKEDSAEPA  
 PVVEVGGEVESKPEEKVAVKPESQPSDKPAEESKVEQAGEPVAPREDEKAPVEPEKQPEAPEEEKAVEE  
 TPKQEESTPDTKAEETVEPKEETVNQSIQPKVETPAVEKQTEPTEEPKVEQAGEPVAPREDEQAPTAP  
 VEPEKQPEVPPEEKA VEETPKPEDKIKGIGTKPEVDKSELNNQIDKASSVSPTDYSTASYNALGPVLET  
 AKGVYASEPVKQPEVNSETNKLKTAIDALNVDKTELNNTIADAKTKVKEHYSDRSWQNLQTEVTKAEKV  
 AANTDAKQSEVNEAVEKLATIEKLVELSEKPIILTLTSTDKKILEREAVAKYTLNQNKTIKSITAEL  
 KKGEVINTVVLTDKVTETETISAAFNLEYKEYTLSTMIYDRNGEETETLENQNIQLDLKKVELK  
 NIKRTDLIKYENGKETNESLITTIIPDDKSNNYLKITSNNQKTLLAVKNIEETTNGTPVYKVTAIADN  
 LVSRTADNKFEE

**SP124 amino acid (SEQ ID NO:219)**

AACACCTGTATATAAAGTTACAGCAATCGCAGACAATTTAGTCTCTAGAACTGCTGATAATAAATTTGA  
 AGAAGAATACGTTCACTATATTGAAAAACCTAAAGTCCACGAAGATAATGTATATTATAATTTCAAAGA  
 ATTAGTGGAAGCTATTCAAACGATCCTTCAAAGAATATCGTCTGGGACAATCAATGAGCGCTAGAAA  
 TGTTGTTCTTAATGGAAGATCATATATCACTAAAGAATTCACAGGAAAACTTTTAAAGTTCTGAAGGAAA  
 ACAATTTGCTATTACTGAATTGGAACATCCATTATTTAATGTGATAACAAACGCAACGATAAATAATGT

Table 1

94

GAATTTTGAAAATGTAGAGATAGAACGTTCTGGTCAAGATAATATTGCATCATTAGCCAATACTATGAA  
 AGGTTCTTCAGTTATTACAAATGTCAAATTTACAGGCACACTTTCAGGTCGTAATAATGTTGCTGGATT  
 TGTAATAATATGAATGATGGAACCTCGTATTGAAAATGTTGCTTTCTTTGGCAAACCTACACTCTACAAG  
 TGGAAATGGCTCTCATACAGGGGGAATTGCAGGTACAACTATAGAGGAATTGTTAGAAAAGCATATGT  
 TGATGCTACTATTACAGGAAACAAAACACGCCAGCTTGTAGTTCTTAAAGTAGATTATGGATTAACT  
 TCTAGACCATCTTATTGGTACAAAAGCTCTCCTAACTGAGTCGGTTGTAAAAGGTAAAATAGATGTTTC  
 AAATCCAGTAGAAGTTGGAGCAATAGCAAGTAAGACTTGGCCTGTAGGTACGGTAAGTAATTCTGTCAG  
 CTATGCTAAGATTATCCGTGGAGAGGAGTTATTTCGGCTCTAACGACGTTGATGATTCTGATTATGCTAG  
 TGCTCATATAAAAAGATTTATATGCGGTAGAGGGATATTTCGTACGTAATAGATCATTTAGGAAATCTAA  
 AACATTTACTAAATTAATAAGAAACAAGCTGATGCTAAAGTTACTACTTTCAATATTACTGCTGATAA  
 ATTAGAAAGTGATCTATCTCTCTTGCAAAACCTTAATGAAGAAAAAGCCTATTCTAGTATTCAAGATTA  
 TAACGCTGAATATAACCAAGCCTATAAAAATCTTGAAAAATTAATACCATTCTACAATAAAGATTATAT  
 TGTATATCAAGGTAATAAATTAAATAAAGAACACCATCTAAATACTAAAGAAGTTCTTTCTGTTACCGC  
 GATGAACAACAATGAGTTTATCACAACCTAGATGAAGCTAATAAAATTATTGTTCACTATGCGGACGG  
 TACAAAAGATTACTTTAACTTGTCTTCTAGCAGTGAAGGTTTAAGTAATGTAAAAGAATATACTATAAC  
 TGACTTAGGAATTAATATACACCTAATATCGTTCAAAAAGATAACACTACTCTTGTTAATGATATAAA  
 ATCTATTTTAGAATCAGTAGAGCTTCAGTCTCAAACGATGTATCAGCATCTAAATCGATTAGGTGACTA  
 TAGAGTTAATGCAATCAAAGATTTATATTTAGAAGAAAGCTTCACAGATGTTAAAGAAAACCTTAACAAA  
 CCTAATCACAAAATTAGTTCAAAACGAAGAACATCAACTAAATGATTCTCCAGCTGCTCGTCAAATGAT  
 TCGTGATAAAGTCGAGAAAAACAAAGCAGCTTTATTACTAGGTTTAACTTACCTAAATCGTTACTATGG  
 AGTTAAATTTGGTGATGTTAATATTAAAGAATTAATGCTATTCAAAACCAGATTTCTATGGTGAAAAAGT  
 TAGCGTATTAGACAGATTAATTGAAATCGGTTCTAAAGAGAACAAACATTAAAGGTTACGTACATTCTGA  
 CGCATTCCGTCAAGTA

**SP124 amino acid (SEQ ID NO:220)**

TPVYKVTAIADNLVSRTADNKFEEYVHYIEKPKVHEDNVYYNFKELVEAIQNDPSKEYRLGQSMSARN  
 VVPNGKSYITKEFTGKLLSSEKQFAITELEHPLFNVTNATINNPNFENVEIERSGQDNIAASLANTMK  
 GSSVITNVKITGTLSGRNNVAGFVNNDGTRIENVAFGKLHSTSGNGSHTGGIAGTNYRGIVRKAYV  
 DATITGNKTRASLLVPKVDYGLTLDHLIGTKALLTESVVKGKIDVSNPVEVGAIASKTWPVGTVSNSVS  
 YAKIIRGEELFGSNDVDDSDYASAHIKDLYAVEGYSSGNRSFRKSKTFTKLTKQADAKVTTFNITADK  
 LESDLSPLAKLNEEKAYSSIQDYNAEYNQAYKNLEKLIPFYNKDYIVYQGNKLNKEHHLNLTKEVLSVTA  
 MNNNEFITNLDEANKIIVHYADGTDYFNLSSSSEGLSNVKEYTITDLGIKYTPNIVQKDNTTLVNDIK  
 SILESVELQSOTMYQHLNRLGDYRVNAIKDLYLEESFTDVKENLTNLITKLQNEEHQLNDSPAARQMI  
 RDKVEKNKAALLLGLTYLNRYYGVKFGDVNIKELMLFKPDFYGEKVSVDRLIEIGSKENNIKGSRTFD  
 AFGQV

**SP125 nucleotide (SEQ ID NO:221)**

ATTAGACAGATTAATTGAAATCGGTTCTAAAGAGAACAAACATTAAAGGTTACGTACATTCGACGCATT  
 CGGTCAAGTATTGGCTAAATATACTAAATCAGGTAATTTAGATGCATTTTTAAATTATAATAGACAATT  
 GTTCACAAATATAGACAATATGAACGATTGGTTTATTGATGCTACAGAAGACCATGTCTACATCGCAGA  
 ACGCGCTTCTGAGGTCGAAGAAATTAATAATCTAAACATCGTGCAATTCGATAATTTAAACGAAGTCA  
 CCTTAGAAATACTATACTCCCACTACTGAATATTGATAAAGCACATCTTTATTTAATTTCAAATTATAA  
 TGCAATTGCCTTTGGTAGTGCAGAGCGATTAGGTAAAAATCATTAGAAGATATTAAAGATATCGTTAA  
 CAAAGCTGCAGATGGTTATAGAAACTATTATGATTTCTGGTATCGTCTAGCGTCTGATAACGTTAAACA  
 ACGACTACTAAGAGATGCTGTTATTCCTATTTGGGAAGGTTATAACGCTCCTGGTGATGGGTTGAAAA  
 ATATGGCCGCTATAATACCGACAAAGTATATACTCCTCTTAGAGAATTCTTTGGTCCTATGGATAAGTA  
 TTATAATTATAATGGAACAGGAGCTTATGCTGCTATATATCCTAACTCTGATGATATTAGAAGTATGATG  
 AAAATATGTTTCATTTAGAAATGGTTGGTGAATACGGTATTTTCAGTTTACACACATGAAACAACACACGT  
 CAACGACCGTGCGATTTACTTAGGTGGCTTTGGACACCGTGAAGGTAAGTATGCTGAAGCATATGCTCA  
 GGGTATGCTACAACTCCTGTTACTGGTAGTGGATTTGATGAGTTTGGTCTTTAGGTATTAATATGGT  
 ATTTAAACGCAAAATGATGGGAATCAGTGGTATATTACAGATCCAAAACTCTAAAAACACGAGAAGA  
 TATTAATAGATATATGAAGGGTTATAATGACACTTTAACTCTTCTTGATGAAATTGAGGCTGAATCTGT  
 GATTTCTCAACAAAATAAAGATTTAAATAGTGCATGGTTCAAAAAAATAGATAGAGAATACCGTGATAA  
 CAATAAATTAATCAATGGGATAAAATTCGAAATCTAAGTCAAGAAGAGAAAAATGAATTAATATTCA  
 ATCTGTTAATGATTTAGTTGATCAACAATTAATGACTAATCGCAATCCAGGTAATGGTATCTATAAACC  
 CGAAGCAATTAGCTATAACGATCAATCACCTTATGTAGGTGTTAGAATGATGACCGGTATCTACGGAGG  
 TAATACTAGTAAAGGTGCTCCTGGAGCTGTTTCATTCAAACATAATGCTTTTAGATTATGGGGTTACTA  
 CGGATACGAAAATGGGTTCTTAGGTTATGCTTCAATAAATAAACAACAATCTAAACAGATGGTGA

Table 1

GTCTGTTCTAAGTGATGAATATATTATCAAGAAAATATCTAACAATACATTTAATACTATTGAAGAATT  
TAAAAAAGCTTACTTCAAAGAAGTTAAAGATAAAGCAACGAAAGGATTAAACAACATTCGAAGTAAATGG  
TTCTTCCGTTTCATCATACGATGATTTACTGACATTGTTTAAAGAAGCTGTTAAAAAAGATGCCGAAAC  
TCTTAAACAAGAAGCAAACGGTAATAAAACAGTATCTATGAATAATACAGTTAAATTAAAAGAAGCTGT  
TTATAAGAACTTCTTCAACAAACAAATAGCTTTAAACTTCAATCTTTAAA

**SP125 amino acid (SEQ ID NO:222)**

LDRLIEIGSKENNIKGSRTFDAGQVLAKYTKSGNLDAFLNYNRQLFTNIDNMNDWFDATEDHVIYIAE  
RASEVEEIKNSKHAFDNLKRSHLRNTILPLLNIDKAHLYLISNYNAIAFGSAERLGKKSLEDIKDIVN  
KAADGYRNYDFWYRLASDNVQRLLRDAVIPIWEGYNAPGWVEKYGRYNTDKVYTPLREFFGPMCKY  
YNYNGTGAYAAIYPNSDDIRTDVKYVHLEMVGEYGISVYTHETHVNDRAIYLGFGHREGTDAEAYAQ  
GMLQTPVTGSGFDEFGSLGINMVFKRKNQWYITDPKTLKTREDINRYMKGYNLTLTLLDEIEAESV  
ISQONKDLNSAWFKKIDREYRDNNKLNQWDKIRNLSQEEKNELNIQSVNDLVDQQLMTRNPNPNGIYKP  
EASYNQSPYVGVRMMTGIYGGNTSKGAPGAVSFKHNAFRLWGYGYENGFLGYASNKYKQSKTDGE  
SVLSDEYIIKKISNNTFNTIEEFKKAYFKEVKDKATKGLTTFEVNGSSVSSYDDLTLFKEAVKKDAET  
LKQEANGNKTVSMNNTVKLKEAVYKLLQQTNSFKTSIFK

**SP126 nucleotide (SEQ ID NO:223)**

TAAGACAGATGAACGGAGCAAGGTGTTTGACTTTTCCATTCCCTACTATACTGCAAAAAATAAACTCAT  
TGTCAAAAAATCTGACTTGACTACTTATCAGTCTGTAAACGACTTGGCGCAGAAAAAGGTTGGAGCGCA  
GAAAGGTTTCGATTCAAGAGACGATGGCGAAAGATTTGCTACAAAAATCTTCCCTCGTATCTCTGCCTAA  
AAATGGGAATTTAATCACAGATTTAAAATCAGGACAAGTGGATGCCGTTATCTTTGAAGAACCTGTTTC  
CAAGGGATTTGTGGAAAATAATCCTGATTTAGCAATCGCAGACCTCAATTTTGAAAAAGAGCAAGATGA  
TTCTACGCGGTAGCCATGAAAAAGATAGCAAGAAATTGAAGAGGCAGTTCGATAAAACCATTCAAAA  
GTTGAAGGAGTCTGGGGAATTAGACAAACTCATTGAGGAAGCCTTA

**SP126 amino acid (SEQ ID NO:224)**

KTDERSKVDFDSIPYYTAKNKLIVKSDLTTYQSVNDLAQKKVGAQKGSIQETMAKDLLQNSSLVSLPK  
NGNLITDLKSGQVDAVIFEPPVSKGFVENNPDLAIADLNFEKEQDDSYAVAMKKDSKKLKRQFDKTIQK  
LKESGELDKLIEEAL

**SP127 nucleotide (SEQ ID NO:225)**

CTGTGAGAATCAAGCTACACCCAAAGAGACTAGCGCTCAAAAGACAATCGTCCTTGCTACAGCTGGCGA  
CGTGCCACCATTGACTACGAAGACAAGGGCAATCTGACAGGCTTGATATCGAAGTTTAAAGGCAGT  
AGATGAAAAACTCAGCGACTACGAGATTCAATTCCAAAGAACCGCTGGGAGAGCATCTTCCAGGACT  
TGATTCTGGTCACTATCAGGCTGCGGCCAATAACTTGAGTTACACAAAAGAGCGTGCTGAAAAATACCT  
TACTCGCTTCCAATTTCCAACAATCCCTCGTCTTGTGAGCAACAAGAAAAATCCTTTGACTTCTCT  
TGACCAGATCGCTGGTAAAACAACACAAGAGGATACCGGAACCTTCTAACGCTCAATTCATCAATAACTG  
GAATCAGAAACACACTGATAATCCCGCTACAATTAATTTTCTGGTGAGGATATTGGTAAACGAATCCT  
AGACCTTGCTAACGGAGAGTTTGATTTCCTAGTTTTTGACAAGGTATCCGTTCAAAGATTATCAAGGA  
CCGTGGTTTAGACCTCTCAGTCGTTGATTTACCTTCTGCAGATAGCCCCAGCAATTATATCATTTCCTC  
AAGCGACCAAAAGAGTTTAAAGAGCAATTTGATAAAGCGCTCAAAGAACTCTATCAAGACGGAACCTT  
TGAAAAACTCAGCAATACCTATCTAGGTGGTTCTTACCTCCAGATCAATCTCAGTTACAA

**SP127 amino acid (SEQ ID NO:226)**

CENQATPKETSAQKTIVLATAGDVPPFDYEDKGNLTGFDIEVLKAVDEKLSDYEIQFORTAWESIFPGL  
DSGHYQAAANNLSYTKERAELYSLPISNNPLVLVSNKKNPLTSLDQIAGKTTQEDTGTSNAQFINNW  
NQKHTDNPATINFSGEDIGKRILDLANGEFDLFLVDKVSQKIIKDRGLDLSVVDLPSADSPSNYIIFS  
SDQKEFKEQFDKALKELYQDGTLEKLSNTYLGGSYLPDQSQLQ

Table 2

*S. pneumoniae* Antigenic Epitopes**SP001**

Lys-1 to Ile-10; Leu-13 to Lys-32; Arg-41 to Ile-51; Ser-85 to Glu-97; Ala-159 to His-168; Val-309 to Thr-318; Val-341 to Asn-352; Asn-415 to Met-430; Phe-454 to Asn-464; Ser-573 to Gly-591; Asn-597 to Thr-641; and Asn-644 to Ala-664.

**SP004**

Thr-9 to Thr-24; Ile-29 to Ala-48; Thr-49 to Val-56; Val-286 to Val-312; Pro-316 to Glu-344; Val-345 to Ile-367; Gln-368 to Val-399; Ser-400 to Glu-431; Asn-436 to Ala-457; Ile-467 to Ala-498; and Thr-499 to Glu-540.

**SP006**

Glu-1 to Lys-13; Pro-24 to Gly-36; Val-104 to Thr-112; Ala-118 to Asn-130; Trp-137 to Ala-146; Ser-151 to Ile-159; Ile-181 to Leu-188; and Pro-194 to Tyr-202.

**SP007**

Gly-1 to Asn-7; Tyr-24 to Gln-34; His-47 to Phe-55; Ser-60 to Ala-67; Ala-122 to Leu-129; Leu-221 to Lys-230; Val-236 to Phe-256; and Asp-271 to Gly-283; and Leu-291 to Asp-297.

**SP008**

Leu-4 to Lys-17; Gln-24 to Leu-32; Asp-60 to Ser-66; Ser-70 to Asp-76; Ala-276 to Lys-283; Asn-304 to Lys-311; and Thr-429 to Pro-437.

**SP009**

Thr-4 to Glu-11; Leu-50 to Asp-60; Ile-102 to Trp-123; and Ser-138 to Ile-157.

**SP010**

Phe-34 to Gly-41; Asp-44 to Lys-50; Leu-172 to Val-186; Leu-191 to Val-198; Ser-202 to Ile-209; and Val-213 to Leu-221.

**SP011**

Asn-2 to Thr-10; Asp-87 to Ala-102; Tyr-125 to Glu-132; Thr-181 to Tyr-189; Arg-217 to Thr-232; Asn-257 to Lys-264; Pro-271 to Ser-278; Tyr-317 to Ala-325; Glu-327 to Pro-337; and Thr-374 to Val-381.

**SP012**

Gly-1 to Lys-19; Phe-34 to Tyr-41; Leu-109 to Lys-126; and Leu-231 to Glu-247.

**SP013**

Ala-1 to Lys-12; Ile-42 to Pro-53; Leu-138 to Lys-146; Ile-205 to Lys-217; Ser-235 to Ile-251; and Ser-261 to Tyr-272.

**SP014**

Gly-1 to Val-16; Leu-35 to Leu-44; Asp-73 to Asp-81; Ile-83 to Asp-92; Glu-145 to Ile-153; Phe-188 to Asn-196; Ser-208 to Phe-215; Ile-224 to Leu-231; and Asn-235 to Ala-243.

**SP015**

Ser-1 to Pro-16; Asn-78 to Glu-88; Ala-100 to Val-108; Ala-122 to Thr-129; Thr-131 to Ser-137; Leu-201 to Ser-220; and Gly-242 to Val-251.

Table 2

*S. pneumoniae* Antigenic Epitopes**SP016**

Gly-1 to Glu-20; Thr-30 to Val-38; Gln-94 to Asn-105; Lys-173 to Pro-182; Gly-189 to Arg-197; Ser-207 to Val-224; Pro-288 to Leu-298; Ala-327 to Ala-342; and Ser-391 to Ala-402.

**SP017**

Ser-1 to Thr-12; Ala-36 to Tyr-45; Gln-48 to Ile-54; Lys-59 to Lys-76; Tyr-113 to Leu-138; and Phe-212 to Asp-219.

**SP019**

Val-97 to Glu-117; Asp-163 to Leu-169; Thr-182 to Thr-191; and Lys-241 to Ser-250.

**SP020**

Asn-18 to Lys-25; Thr-47 to Glu-60; Trp-75 to Val-84; Gly-102 to Val-110; Pro-122 to Ala-131; and Glu-250 to Pro-258.

**SP021**

Ser-1 to Asp-8; Val-44 to Asp-54; Ala-117 to Val-125; Thr-165 to Thr-173; and Glu-180 to Pro-189.

**SP022**

Phe-5 to Lys-13; Thr-20 to Ser-36; Glu-59 to Lys-81; Tyr-85 to Gly-93; Trp-94 to Trp-101; and Thr-195 to Trp-208.

**SP023**

Gln-45 to Glu-59; Asp-69 to Pro-85; Lys-111 to Asn-121; Pro-218 to Ala-228; and Glu-250 to Asn-281.

**SP025**

Gln-14 to Thr-20; Gly-27 to Phe-33; Gly-63 to Glu-71; and Ile-93 to Phe-102.

**SP028**

Asp-171 to Pro-179; Tyr-340 to Glu-350; Pro-455 to Tyr-463; and Asp-474 to Pro-480.

**SP030**

Leu-22 to Leu-37; Trp-81 to Ala-90; Phe-101 to Ala-106; Thr-124 to Tyr-130; and Asn-138 to Glu-144.

**SP031**

Asp-8 to Val-16; Gly-27 to Thr-35; Gly-178 to Asp-195; Thr-200 to Asp-209; Trp-218 to Leu-224; and Lys-226 to Asp-241.

**SP032**

Ser-9 to Asp-28; Phe-31 to Val-40; Gly-42 to Arg-50; Ile-52 to Leu-60; Asp-174 to Phe-186; Leu-324 to Met-333; and Thr-340 to Asn-347.

**SP033**

Gln-2 to Ile-13; Phe-46 to Ile-53; and Asp-104 to Thr-121.

**SP034**

Glu-36 to Gly-43; Ala-188 to Asp-196; Trp-313 to Gly-320; and Leu-323 to Leu-329.

Table 2

*S. pneumoniae* Antigenic Epitopes**SP035**

Arg-19 to Asp-36; Asp-47 to Val-57; Asn-134 to Thr-143; Asp-187 to Arg-196; and Glu-222 to Ser-230.

**SP036**

Arg-10 to Arg-17; Lys-29 to Ser-39; Ser-140 to Ala-153; Arg-158 to Tyr-169; Asp-175 to Ala-183; Gly-216 to Asn-236; Ala-261 to Leu-270; Arg-282 to Phe-291; and Thr-297 to Ala-305; Pro-342 to Gln-362; Phe-455 to Asp-463; His-497 to Thr-511; Ala-521 to Gly-529; Ile-537 to Val-546; Ile-556 to Ala-568; Pro-581 to Ser-595; Glu-670 to Ala-685; Ser-696 to Ala-705 and Leu-782 to Ser-791.

**SP038**

Glu-61 to Pro-69; Phe-107 to Ala-115; Leu-130 to Tyr-141; Ala-229 to Glu-237; Ser-282 to Asn-287; Ala-330 to Glu-338; and Tyr-387 to Glu-393.

**SP039**

Ser-28 to Asp-35; Pro-88 to Pro-96; Leu-125 to Arg-135; Phe-149 to Leu-157; Gln-246 to Val-254; Ala-357 to Thr-362; Gly-402 to Lys-411; and Leu-440 to Pro-448.

**SP040**

Thr-21 to Ile-30; His-54 to Gln-68; Arg-103 to Leu-117; and Thr-127 to Leu-136.

**SP041**

Gly-36 to Asp-49; Leu-121 to Val-128; and Ala-186 to Ile-196.

**SP042**

Gly-11 to Arg-19; Ile-23 to Lys-31; His-145 to Asn-151; Gln-159 to Asp-166; Ile-175 to Asp-181; Gly-213 to Tyr-225; Ile-283 to Val-291; Pro-329 to Glu-364; Arg-372 to Ser-386; Thr-421 to Phe-430; Leu-445 to Val-453; Ile-486 to Ala-497; Asp-524 to Ala-535; His-662 to Gly-674; and His-679 to Gln-702.

**SP043**

Lys-2 to Asp-12; Val-58 to Asn-68; Ser-87 to Asp-95; and Asp-102 to Lys-117.

**SP044**

Gln-3 to Lys-11; Asp-37 to Tyr-52; Glu-171 to Leu-191; His-234 to Asn-247; and Asn-283 to Ala-291.

**SP045**

Tyr-52 to Ile-63; Asp-212 to Gln-227; Ser-315 to Thr-332; Leu-345 to Phe-354; Asp-362 to Val-370; Thr-518 to Asn-539; Ala-545 to Lys-559; and Val-601 to Pro-610.

**SP046**

Gln-9 to Ala-18; Glu-179 to Lys-186; Lys-264 to Glu-271; Gly-304 to Glu-17; Ser-503 to Asn-511; Asn-546 to Thr-553; and Asn-584 to Asp-591.

**SP048**

Table 2

*S. pneumoniae* Antigenic Epitopes

Tyr-4 to Asp-25; Lys-33 to Val-70; Asp-151 to Thr-170; Asp-222 to Val-257; Thr-290 to Phe-301; and Gly-357 to Val-367.

**SP049**

Ala-23 to Arg-37; Tyr-85 to Gln-95; Glu-106 to Ile-118; Arg-131 to Ile-144; Gly-150 to Ser-162; and Ala-209 to Asp-218.

**SP050**

Asp-95 to Glu-113; Gly-220 to Gly-228; Asn-284 to Glu-295; Thr-298 to Val-315.

**SP051**

Lys-16 to Glu-50; Lys-57 to Asn-104; Ser-158 to Trp-173; Asp-265 to Pro-279; Val-368 to Tyr-386; Glu-420 to Ile-454; Pro-476 to Ile-516; Phe-561 to Gly-581; Thr-606 to Gly-664; and Glu-676 to Val-696.

**SP052**

Asn-41 to Tyr-60; Phe-80 to Glu-103; Ala-117 to Val-139; Ile-142 to Leu-155; Val-190 to Lys-212; Glu-276 to Phe-283; Arg-290 to Ser-299; Leu-328 to Val-351; Gly-358 to Thr-388; Glu-472 to Ala-483; Val-533 to Asn-561; Asp-595 to Val-606; Glu-609 to Val-620; Glu-672 to Ser-691.

**SP053**

Ala-62 to Val-101; Thr-147 to Leu-174; Lys-204 to Val-216; Gln-228 to Val-262; Ser-277 to Gly-297; Thr-341 to Gly-368; Thr-385 to Ala-409; Thr-414 to Ser-453; Asn-461 to Leu-490; Glu-576 to Thr-625; Gly-630 to Arg-639; and Asp-720 to Leu-740.

**SP054**

Glu-7 to Val-28; and Tyr-33 to Glu-44.

**SP055**

Pro-3 to Val-18; Thr-21 to Lys-53; Val-84 to Lys-99; Ile-162 to Val-172; and Val-204 to Ser-241.

**SP056**

Val-34 to Tyr-41; Leu-47 to Glu-55; and Pro-57 to Gln-66.

**SP057**

Asp-1 to Val-25; Pro-29 to Ile-80; Asn-96 to Val-145; and Pro-150 to Glu-172.

**SP058**

Ala-64 to Thr-70; Leu-82 to His-138; and Val-228 to Asn-236.

**SP059**

Val-10 to Thr-24; Ser-76 to Pro-102; Ser-109 to Ile-119; Ser-124 to Val-130; Thr-186 to Ile-194; and Asn-234 to Ser-243.

**SP060**

Leu-70 to Arg-76; and Val-79 to Ile-88.

**SP062**

Glu-14 to Lys-28; Ser-32 to Lys-46; and Glu-66 to Thr-74.

Table 2

*S. pneumoniae* Antigenic Epitopes**SP063**

Ile-10 to Val-25; Val-30 to Thr-40; Asp-44 to Pro-54; Asn-57 to Val-63; Pro-71 to Val-100; and Thr-105 to Thr-116.

**SP064**

Pro-12 to Leu-32; Val-40 to Leu-68; Asp-95 to Ala-125; Ser-164 to Glu-184; Ser-314 to Glu-346; Asn-382 to Val-393; Leu-463 to Gln-498; Asn-534 to Lys-548; and Lys-557 to Gly-605.

**SP065**

Asn-2 to Ile-12; Ala-39 to Thr-61; and His-135 to Ala-155.

**SP067**

Gly-1 to Thr-13; Asp-203 to Asn-218; and Gly-240 to Asp-253.

**SP068**

Ser-2 to Ser-12; Val-17 to Gln-26; and Lys-54 to Cys-67.

**SP069**

Ser-32 to Thr-41; Pro-66 to Glu-80; Thr-110 to Val-122; and Val-147 to Thr-180.

**SP070**

Lys-6 to Tyr-16; Gln-19 to Ile-27; Arg-50 to Ala-58; Leu-112 to Val-128; Ile-151 to Asn-167; Leu-305 to Phe-321.

**SP071**

Gln-92 to Asn-158; Gln-171 to Gln-188; Val-204 to Val-240; Thr-247 to Ala-273; Glu-279 to Thr-338; Pro-345 to Glu-368; Asn-483 to Lys-539; Val-552 to Ala-568; Glu-575 to Ser-591; Ser-621 to Gly-640; Gln-742 to Gly-758.

**SP072**

Val-68 to Tyr-81; Tyr-86 to Val-121; Leu-127 to Gly-140; Gly-144 to Ala-155; Gln-168 to Val-185; Asp-210 to Try-241; Glu-246 to Thr-269; Lys-275 to Tyr-295; Gly-303 to Pro-320; Arg-327 to Ile-335; Thr-338 to Thr-364; Tyr-478 to Phe-495; and Tyr-499 to Arg-521.

**SP073**

Glu-37 to Val-45; Glu-55 to Val-68; Thr-104 to Thr-119; Ile-127 to Tyr-135; Asn-220 to Ile-232; Thr-237 to Ala-250; Ser-253 to Ala-263; Glu-284 to Ile-297; and Met-438 to Asn-455.

**SP074**

Gly-2 to Ala-12; Gly-96 to Ile-110; and Thr-220 to Phe-239.

**SP075**

Phe-33 to Tyr-42; Gln-93 to Gly-102; and Val-196 to Asp-211.

**SP076**

Ser-64 to Leu-76; and Phe-81 to Ala-101.

**SP077**

Asp-1 to Glu-12; Tyr-26 to Val-36; and Val-51 to Try-62.



Table 2

*S. pneumoniae* Antigenic Epitopes**SP078**

Ala-193 to Ile-208; Tyr-266 to Asn-275; Glu-356 to Leu-369; Ala-411 to Gly-422; Ser-437 to Pro-464; Thr-492 to Glu-534; and Glu-571 to Gln-508.

**SP079**

Gly-11 to Leu-20; Lys-39 to Leu-48; Leu-72 to Val-85; Asn-147 to Ser-158; Ile-178 to Asp-187; Tyr-189 to Gln-201; and Leu-203 to Ala-216

**SP080**

Ser-2 to Glu-12; Gln-42 to Ala-51; Ala-116 to Ser-127; Phe-131 to Asp-143; and Ile-159 to Ile-171.

**SP081**

Gln-2 to Leu-9; Gln-49 to Cys-57; Ile-108 to Val-131; Gly-134 to Leu-145; and Trp-154 to Cys-162.

**SP082**

Ile-101 to Ser-187; Gly-191 to Asn-221; Arg-225 to Arg-236; Tyr-239 to Leu-255; and Gly-259 to Arg-268.

**SP083**

Ser-28 to Asp-70.

**SP084**

Leu-42 to Gln-66; Thr-69 to Lys-81; Glu-83 to Arg-92; and Gly-98 to Asn-110.

**SP085**

Gln-2 to Val-22; and Ser-45 to Glu-51.

**SP086**

Leu-18 to Gln-65; and Lys-72 to Val-83.

**SP087**

Ser-45 to Leu-53; and Thr-55 to Gln-63

**SP088**

Pro-8 to Ile-16; Leu-25 to Trp-33; Tyr-35 to Gln-43; Leu-51 to Val-59; Val-59 to Arg-67; Thr-55 to Tyr-63; Asn-85 to Gly-93; Thr-107 to Leu-115; Leu-115 to Trp-123; Ala-121 to Thr-129; Tyr-153 to Ala-161; His-176 to Gly-184; Tyr-194 to Ala-202; Ala-217 to Gly-225; and Asn-85 to Gly-93.

**SP089**

Trp-43 to Ala-51; Gln-68 to Phe-76; Val-93 to Gln-101; Phe-106 to Phe-114; Lys-117 to Lys-125; Trp-148 to Phe-156; Glu-168 to Gln-176; Ile-193 to Tyr-201; Lys-203 to Lys-211; Glu-212 to Gln-220; Ile-237 to Tyr-245; Lys-247 to Lys-255; Glu-256 to Gln-264; Met-275 to Gly-283; Lys-286 to Gly-294; Trp-292 to Glu-300; Asp-289 to Thr-297; Tyr-315 to Ser-323; Asp-334 to Lys-342; Pro-371 to Arg-379; Arg-485 to Asn-493; Lys-527 to Arg-535; Phe-537 to Met-545; and Tyr-549 to Glu-557.

**SP090**

Table 2

*S. pneumoniae* Antigenic Epitopes

Phe-2 to Gln-10; Gln-13 to Lys-21; Tyr-19 to Glu-27; Tyr-39 to Met-47; Pro-65 to Leu-73; Tyr-121 to His-129; Lys-147 to Ile-155; Gly-161 to Lys-169; Gly-218 to Trp-226; Asp-230 to Thr-238; Tyr-249 to Ala-257; and Ala-272 to Gly-280.

**SP091**

Ser-19 to Ser-27; Asn-25 to Thr-33; Val-51 to Gln-59; Asn-75 to Asn-83; Ile-103 to Trp-111; Tyr-113 to Ala-121; Leu-175 to Asn-183; Glu-185 to Trp-193; Ala-203 to Tyr-211; Val-250 to Phe-258; Asn-260 to Thr-268; Ser-278 to Asp-286; Tyr-305 to Leu-313; Asn-316 to Gly-324; Asn-374 to Asp-382; Asn-441 to Gly-449; and Ser-454 to Gln-462.

**SP092**

Arg-95 to Glu-103; Ala-216 to Val-224; Leu-338 to Glu-346; Pro-350 to Ala-358; Pro-359 to Ala-367; Pro-368 to Ala-376; Pro-377 to Ala-385; Pro-386 to Ala-394; Pro-395 to Ala-403; Pro-350 to Ala-358; Gln-414 to Lys-422; Pro-421 to Asn-429; Trp-465 to Tyr-473; Phe-487 to Tyr-495; Asn-517 to Gly-525; Trp-586 to Tyr-594; Phe-608 to Tyr-616; and Asp-630 to Gly-638.

**SP093**

Gln-30 to Ile-38; Gln-52 to Val-60; Ala-108 to His-116; Tyr-133 to Glu-141; Tyr-192 to Ala-200; and Phe-207 to Ser-215.

**SP094**

Ala-87 to Val-95; Leu-110 to Cys-118; Gln-133 to Leu-141; Ser-185 to Leu-193; Ile-195 to Gly-203; Asp-206 to Gln-214; Ser-211 to Gly-219; Ile-241 to Thr-249.

**SP095**

Arg-1 to Gln-9; Phe-7 to Asn-15; Thr-21 to Asn-30; Leu-46 to Phe-54; and Ser-72 to Met-80.

**SP096**

Gly-29 to Ile-37; Glu-52 to Ser-60; and Leu-64 to Gly-72.

**SP097**

Ala-11 to Thr-19; Glu-53 to Glu-61; Ser-91 to Lys-99; Thr-123 to Gln-131; and Gly-209 to Lys-217.

**SP098**

Thr-3 to Ser-11; Gly-38 to Phe-46; Tyr-175 to Asn-183; Met-187 to Cys-195; Gln-197 to Leu-205; Tyr-307 to Gln-315; Gly-318 to Tyr-326; Asn-348 to Val-356; Lys-377 to Pro-385; and Leu-415 to Val-423.

**SP099**

Arg-19 to Gly-27; Asp-76 to Ser-84; Val-90 to Lys-98; Phe-165 to Val-173; Leu-237 to Pro-245.

**SP100**

His-111 to Gln-119; Ser-141 to His-149; Asp-154 to Ser-162; Gln-158 to Gln-166; Asp-154 to Gln-166; Lys-180 to Gln-188; and Ser-206 to Gln-214.

**SP101**

Table 2

***S. pneumoniae* Antigenic Epitopes**

Glu-23 to Glu-31; Glu-40 to Val-48; Gln-50 to Ser-58; Thr-61 to Ile-69; Leu-82 to Ile-90; Ala-108 to Leu-116; Gln-121 to Pro-129; and Leu-130 to Thr-138.

**SP102**

Asp-32 to His-40; Arg-48 to Lys-56; and Asp-102 to Thr-110.

**SP103**

Arg-5 to Gln-13; Gln-22 to Leu-30; Arg-151 to Gln-159; Arg-167 to Gln-175; Pro-189 to Glu-197; Gly-207 to Leu-215; Ser-219 to Gln-227; Ser-233 to Ser-241; Pro-255 to Asp-264; Lys-272 to Gly-280; Ser-318 to Val-326; Thr-341 to Asp-351; Asn-356 to Thr-364; Val-370 to Tyr-378; Ile-379 to Gln-387; and Met-435 to Tyr-443.

**SP105**

Asn-28 to Pro-36; Thr-77 to Phe-85; Arg-88 to Val-96; Gly-107 to Phe-115; Asp-169 to Asp-177; His-248 to Ser-256; and Ser-274 to Ala-282.

**SP106**

Val-10 to Thr-18; Ile-62 to Tyr-70; Ile-71 to Pro-79; Lys-86 to Gln-94; Lys-100 to Thr-108; Phe-132 to Leu-140; and Asp-145 to Arg-153.

**SP107**

Asp-33 to Val-41; and Arg-63 to Gln-71.

**SP108**

Lys-9 to Gln-17; Leu-44 to Ser-52; Ser-63 to Phe-71; Tyr-109 to Ser-117; Ile-183 to Ile-191; Pro-194 to Leu-202; Gly-257 to Gln-265; Ala-323 to Thr-331; and Leu-381 to Tyr-389.

**SP109**

Asn-2 to Gln-10; Ala-65 to Lys-73; Leu-76 to Glu-84; Thr-111 to Asp-119; Gln-116 to Tyr-124; Tyr-130 to Val-138; Asp-173 to Gly-181; Asp-196 to Ser-204; Asn-231 to Ser-239; Phe-252 to Ser-260; Phe-270 to Tyr-278; Val-291 to His-299; Asp-306 to Leu-314; and Pro-327 to Gly-335.

**SP110**

Ser-8 to Glu-16; Ile-37 to Val-45; Ala-107 to Val-115; and Gly-122 to Thr-130.

**SP111**

Asp-19 to Glu-28; Leu-43 to Ala-51; Asn-102 to Phe-110; Gln-133 to Ser-141; Phe-162 to Asp-170; Tyr-194 to Met-202; and Asp-273 to Ser-281.

Table 2

*S. pneumoniae* Antigenic Epitopes**SP112**

Asp-3 to Gln-11; Gly-21 to Ile-29; Ala-46 to Arg-54; Arg-98 to Arg-106; Thr-114 to Val-122; Gln-133 to Asn-141; and Leu-223 to Thr-231.

**SP113**

Asn-19 to Gly-27; Arg-54 to Ser-62; Val-69 to Gln-77; Ser-117 to Asn-125; Gly-164 to Leu-172; Tyr-193 to Ser-201; Cys-303 to Phe-311; His-315 to Ile-323; Arg-341 to Cys-349; Ile-347 to Ser-355; Arg-403 to Phe-411; Gln-484 to Pro-492; Ser-499 to Leu-507; Ile-541 to Thr-549  
Asn-622 to Ile-630; and Glu-645 to Gly-653.

**SP114**

Gly-17 to Leu-25; His-40 to Gln-48; Arg-49 to Arg-57; Ile-65 to Pro-73;  
Asn-101 to Asp-111; Gly-128 to Cys-136; Phe-183 to Thr-191; and Pro-268 to Ile-276.

**SP115**

Met-8 to Ser-16; Tyr-24 to Leu-32; Cys-68 to Leu-76; Ser-100 to Pro-108; Thr-193 to Thr-201; Gly-238 to Pro-250; Thr-280 to Phe-288; Pro-303 to Asn-312; Trp-319 to Leu-328; Leu-335 to Leu-344; Lys-395 to Ala-403; Asn-416 to Gln-424; Tyr-430 to Ser-438; Val-448 to Leu-456; Leu-460 to Thr-468; Pro-502 to Thr-510; Lys-515 to Ile-524; Gln-523 to His-532; Tyr-535 to Thr-543; Ser-559 to Pro-567; Thr-572 to Asn-580;  
Val-594 to Arg-602; Arg-603 to Asn-611; Thr-620 to Trp-628; and Tyr-644 to Arg-653.

**SP117**

Ala-6 to Gly-14; Ile-19 to Thr-27; Thr-99 to Leu-107; Ser-117 to Asp-125; His-131 to Val-139; Ile-193 to Gly-201; and Val-241 to Gln-249.

**SP118**

Ser-8 to Trp-23; His-46 to Ala-54; Asn-93 to Gly-101; Val-100 to Ser-108; Arg-155 to Asp-163; and His-192 to Leu-200.

**SP119**

Tyr-46 to Lys-54; Ser-93 to Ser-101; Trp-108 to Asn-116; Val-121 to Glu-129; and Tyr-131 to Gln-139.

**SP120**

Ala-57 to Lys-65; Leu-68 to Glu-76; Thr-103 to Tyr-116; Tyr-122 to Val-130; His-163 to Gly-173; Asp-188 to Ser-196; Ser-222 to Ser-231; Phe-244 to Ser-252; Pro-262 to Tyr-270; Val-283 to His-291; and Asp-298 to Leu-306.

**SP121**

Ser-3 to Ala-11; Asp-13 to Leu-21; Ser-36 to Val-44; and Gln-136 to Met-144.

**SP122**

Asn-28 to Lys-36; Glu-39 to Thr-50; Val-54 to Lys-62; Asn-106 to Leu-114; Phe-159 to Gly-167; Asn-172 to Arg-180; Glu-199 to Asn-207;

Table 2

***S. pneumoniae* Antigenic Epitopes**

Lys-230 to His-241; Asn-252 to Gly-263; Met-278 to Ala-287; Thr-346 to Asp-354; Lys-362 to Thr-370; Asp-392 to Asn-405; Asp-411 to Ala-424; Gly-434 to Gly-443; Tyr-484 to Glu-492; Ile-511 to Leu-519; Asn-524 to Asp-538; Glu-552 to Ile-567; Val-605 to Lys-613; Phe-697 to Ala-705; Phe-722 to Leu-730; Leu-753 to Leu-761; Asp-787 to Gln-795; Leu-858 to Asn-866; Ala-892 to Thr-901; Gly-903 to Ile-913; Ile-921 to Asn-931; Asn-938 to Pro-951; Gly-960 to Lys-970; Leu-977 to Asp-985; and Leu-988 to Pro-996.

**SP123**

Val-4 to Asn-12; Glu-47 to Leu-55; Lys-89 to Glu-100; Ser-165 to Thr-173; Lys-234 to Val-242; Ser-258 to Ser-266; Glu-284 to Asn-292; Tyr-327 to Leu-335; Tyr-457 to Thr-465; Tyr-493 to Glu-501; Thr-506 to Tyr-514; Lys-517 to Thr-525; Asn-532 to Gly-540; and Arg-556 to Glu-564.

**SP124**

Arg-16 to Glu-24; Gln-52 to Arg-60; Asn-69 to Tyr-77; Glu-121 to Asn-129; Ala-134 to Val-142; Thr-151 to Ala-159; Asn-164 to Glu-172; His-181 to His-189; Thr-210 to Ala-218; Ser-244 to Val-252; Phe-287 to Tyr-297; Ser-312 to Thr-323; His-433 to Tyr-441; Ser-445 to Asn-453; Asn-469 to Thr-477; Asn-501 to Asn-509; Gln-536 to Ala-547; and Gln-608 to Asp-621.

**SP125**

Ser-9 to Asp-21; Ala-28 to Leu-36; Asn-49 to Phe-57; Val-137 to Arg-145; Asn-155 to Leu-163; Glu-183 to Asp-191; Gly-202 to Tyr-210; Pro-221 to Asp-229; Phe-263 to Ala-271; Phe-300 to Gln-308; Asp-313 to Glu-321; Asn-324 to Asp-332; Ile-346 to Asn-354; Asp-362 to Lys-370; Met-402 to Gly-410; Gly-437 to Gly-445; Ser-471 to Glu-483; Gly-529 to Asp-537; Gln-555 to Val-563; and Leu-579 to Lys-587.

**SP126**

Leu-22 to Thr-30; Val-65 to Leu-73; and Thr-75 to Asp-83.

**SP127**

Glu-2 to Ala-12; Asp-28 to Thr-36; Val-105 to Thr-113; Lys-121 to Thr-129; Trp-138 to Pro-146; Ser-152 to Ile-160; Lys-180 to Asp-188; Leu-194 to Asn-202; and Gly-228 to Thr-236.

Table 3

Primer		<i>S. pneumoniae</i> ORF Cloning Primers	
Name	SEQ ID	Sequence	RE
SP001A	NO:227	GACTGGATCCTAAAATCTACGACAATAAAAAATC	Bam HI
SP001B	NO:228	CTGAGTCGACTGGTTGTGCTGGTTGAG	Sal I
SP004A	NO:229	GTCAGGATCCAAATTACAATACGGACTATG	Bam HI
SP004B	NO:230	CAGTGTGCTACTAAGTCTAGGTCGGAAAC	Sal I
SP006A	NO:231	GACTGGATCCTGAGAATCAAGCTACACCCAAAGAG	Bam HI
SP006B	NO:232	AGTCAAGCTTTTGTAACTGAGATTGATCTGG	Hind III
SP007A	NO:233	GACTGGATCCTGGTAACCGCTCTTCTCGTAACGCAGC	Bam HI
SP007B	NO:234	AGTCAAGCTTTTTTCAGGAAGCTTTTACGCTTCC	Hind III
SP008A	NO:235	AGTCAGATCTTGTGGAAATTTGACAGGTAACAGCAAAAAAGCTGC	Bgl II
SP008B	NO:236	ACTGAAGCTTTTTTGTCTTTTCAAGAATTCATCG	Hind III
SP009A	NO:237	GACTGGATCCTGGTCAAGGAAGCTTCTAAAGAC	Bam HI
SP009B	NO:238	AGTCAAGCTTTCACAAATTCGTTGGTGAAGCC	Hind III
SP010A	NO:239	GACTGGATCCTAGCTCAGGTGGAAACGCTGGTTCATCC	Bam HI
SP010B	NO:240	AGTCAAGCTTATCAACTTTTCCACCTTCAACAACC	Hind III
SP011A	NO:241	GTCAAGATCTCTCCAACCTATGGTAAATCTGCGGATGG	Bgl II
SP011B	NO:242	AGTCCTGCAGATCCACATCCGCTTTCATCGGGTTAAAGAAGG	Pst I
SP012A	NO:243	GACTGGATCCTGGGAAAAATTTAGCGAAACTAGTGG	Bam HI
SP012B	NO:244	GTCAGTGCAGCTGTCCTTCTTTACTTCTTTGGTTGC	Pst I
SP013A	NO:245	GACTGGATCCTGCTAGCGGAAAAAAGATACAACCTTCTGG	Bam HI
SP013B	NO:246	CTGAAAGCTTTTTTGCCAACTCTTCAGCAATCTTGTC	Hind III
SP014A	NO:247	GACTAGATCTTGGCTCAAAAAATACAGCTTCAAGTCC	Bgl II
SP014B	NO:248	AGTCCTGCAGGTTTTTGTGCTTGGTATTGGTCG	Pst I
SP015A	NO:249	GACTGGATCCTAGTACAACTCAAGCACTAGTCAGACAGAG	Bam HI
SP015B	NO:250	CAGTCTGCAGTTTCAAAGCTTTTTGTATGTCTTC	Pst I
SP016A	NO:251	GACTGGATCCTGGCAATTTCTGGCGGAAGTAAAGATGC	Bam HI
SP016B	NO:252	AGTCAAGCTTGTTCATAGCTTTTTTGATTGTTTCG	Hind III
SP017A	NO:253	GACTGGATCCTTCACAAGAAAAAACAATAATGAAGATGG	Bam HI
SP017B	NO:254	AGTCAAGCTTATCGACGTAGTCTCCGCCTTC	Hind III
SP019A	NO:255	GACTGGATCCGAAAGGTCTGTGGTCAAATAATCTTACC	Bam HI
SP019B	NO:256	AGTCAAGCTTAGAGTTAATAGGTGCTTGCCAATAGG	Hind III
SP020A	NO:257	GACTGGATCCAAACTCAGAAAAGAAAGCAGACAATGC	Bam HI
SP020B	NO:258	AGTCAAGCTTCCAAACTGGTTGATCCAAACCATCTG	Hind III
SP021A	NO:259	GACTGGATCCTTCGAAAGGGTCAGAAGGTGCAGACC	Bam HI
SP021B	NO:260	AGTCAAGCTTCTGTAGGCTTGGTGTGCCCCAGTTGC	Hind III
SP022A	NO:261	CTGAGGATCCGGGGATGGCAGCTTTTAAAAATC	Bam HI
SP022B	NO:262	CAGTAAGCTTGTTTACCCATTACCATTTACC	Hind III
SP023A	NO:263	CAGTGGATCCAGACGAGCAAAAAATTAAG	Bam HI
SP023B	NO:264	TCAGAAGCTTGTTTACCCATTACCATTT	Hind III
SP025A	NO:265	GACTGGATCCCTGTGGTGAGGAAGAACTAAAAAG	Bam HI
SP025B	NO:266	CTGAGTCGACAATATTCTGTAGGAATGCTTCGAATTTG	Sal I
SP028A	NO:267	CTGAGGATCCGACTTTTAAACAATAAACTATTGAAGAG	Bam HI
SP028B	NO:268	GTCAGTGCAGGTTGTCACTCCAAAAATCACGG	Pst I
SP030A	NO:269	GACTGGATCCCTTTACAGGTAAACAACATAAGTCGG	Bam HI
SP030B	NO:270	CAGTAAGCTTTTCGAAGTTTGGCTCAGAATTG	Hind III
SP031A	NO:271	GACTGGATCCCCAGGCTGATACAAGTATCGCA	Bam HI
SP031B	NO:272	CAGTAAGCTTATCTGCAGTATGGCTAGATGG	Hind III
SP032A	NO:273	GACTGGATCCGCTCTGTATCATTTGAAAACAAAGAAAC	Bam HI
SP032B	NO:274	CAGTCTGCAGTTTTACTGTTGCTGTGCTTGTG	Pst I
SP033A	NO:275	ACTGAGATCTTGGTCAAAAGGAAAGTCAGACAGGAAAGG	Bgl II
SP033B	NO:276	CAGTAAGCTTATTCCTGAGCTTTTTTGATAAAGGTTGCGCA	Hind III
SP034A	NO:277	ACTGGGATCCGAAGGATAGATATATTTTAGCATTTGAGAC	Bam HI
SP034B	NO:278	AGTCAAGCTTCCATGGTATCAAAGGCAAGACTTGG	Hind III
SP035A	NO:279	GTCAGGATCCGGTAGTTAAAGTTGGTATTAACGG	Bam HI
SP035B	NO:280	AGTCAAGCTTGCAATTTTTGCGAAGTATTTCAAGAG	Hind III
SP036A	NO:281	AGTCGGATCCTTCTTACGAGTTGGGACTGTATCAAGC	Bam HI

Table 3

Primer		<i>S. pneumoniae</i> ORF Cloning Primers	
Name	SEQ ID	Sequence	RE
SP036B	NO:282	AGTCAAGCTTGTGTTTATTTTTCCTTACTTACAGATGAAGG	Hind III
SP038A	NO:283	AGTCGGATCCTACTGAGATGCATCATAATCTAGGAGC	Bam HI
SP038B	NO:284	TCAGCTCGAGTTCTTTGACATCTCCATCATAAGTCGC	Xho I
SP039A	NO:285	GACTGGATCCGGTTTGTGAGAAAGTATTTGCAGGGG	Bam HI
SP039B	NO:286	CAGTAAGCTTGGATTTTTTCATGGATGCAATTTTTTTGG	Hind III
SP040A	NO:287	GACTGGATCCGACAACATTTACTATCCATACAGTAGAGTCAGC	Bam HI
SP040B	NO:288	GACTAAGCTTGGCATAAGGTTGCAATTCTGGATTAATTGG	Hind III
SP041A	NO:289	GACTGGATCCGGCTAAGGAAAGAGTGGATG	Bam HI
SP041B	NO:290	GACTAAGCTTTTCATTTTAAATTGACTATGCGCCCG	Hind III
SP042A	NO:291	GACTGGATCCTTGTTCCTATGAACTTGGTCGTCACC	Bam HI
SP042B	NO:292	CATGAAGCTTATCCTGGATTTTCCAAGTAAATCT	Hind III
SP043A	NO:293	GACTGGATCCTTATAAGGGTGAATTAGAAAAAGG	Bam HI
SP043B	NO:294	GACTAAGCTTCTTATTAGGATTGTTAGTAGTTG	Hind III
SP044A	NO:295	GACTGGATCCGAATGTTTCAGGCTCAAGAAAGTTCAGG	Bam HI
SP044B	NO:296	GACTAAGCTTTTCCCCTGATGGAGCAAAGTAATACC	Hind III
SP045A	NO:297	GACTGGATCCCCTTGGGTGTAACCCATATCCAGCTCCTTCC	Bam HI
SP045B	NO:298	GACTGTCGACTTCAGCTTGTATCTGGGGTTGC	Sal I
SP046A	NO:299	GACTGGATCCTAGTGATGGTACTTGGCAAGGAAAACAG	Bam HI
SP046B	NO:300	ACTGCTGCAGATCTTTGCCACCTAGCTTCTCATTG	Pst I
SP048A	NO:301	GTCAGGATCCTGGGATTCAATATGTCAGAGATGATACTAG	Bam HI
SP048B	NO:302	CTAGAAGCTTACGCACCCATTACCATTTATCATTG	Hind III
SP049A	NO:303	GTCAGGATCCGGATAATAGAGAAGCATTAAAAACC	Bam HI
SP049B	NO:304	AGTCAAGCTTGACAAAATCTTGAAACTCCTCTGGTC	Hind III
SP050A	NO:305	GTCAGGATCCAGATTTTGTGCGAGGAGTGTCATACC	Bam HI
SP050B	NO:306	AGTCAAGCTTTCCCTTTTTTACCCTTACGAATCCAGG	Hind III
SP051A	NO:307	GACTGGATCCATCTGTAGTTTATGCGGATGAAACACTTATTAC	Bam HI
SP051B	NO:308	GACTGTCGACGCTTTGGTAGAGATAGAAGTCATG	Sal I
SP052A	NO:309	GACTGGATCCTTACTTTGGTATCGTAGATACAGCCGGC	Bam HI
SP052B	NO:310	AGTCAAGCTTTGTTAATTGCGTACCTTCTAAGCGACC	Hind III
SP053A	NO:311	GACTGGATCCAGCTAAGGTTGCATGGGATGCGATTTCG	Bam HI
SP053B	NO:312	GACTGTCGACCTGGGCTTTATTAGTTTGACTAGC	Sal I
SP054A	NO:313	CAGTGGATCCCTATCACTATGTAAATAAAGAGA	Bam HI
SP054B	NO:314	ACTGAAGCTTTTCTGTCCCTGTTTGAGGCA	Hind III
SP055A	NO:315	CAGTGGATCCTGAGACTCCTCAATCAATAACAAA	Bam HI
SP055B	NO:316	ACGTAAGCTTATAATCAGTAGGAGAACTGAACT	Hind III
SP056A	NO:317	CAGTGGATCCGGATGCTCAAGAACTGCGG	Bam HI
SP056B	NO:318	GACTAAGCTTTTGCCCTCTCATTCTTGCTTCC	Hind III
SP057A	NO:319	CAGTGGATCCCGACAAAGGTGAGACTGAG	Bam HI
SP057B	NO:320	ACGTAAGCTTATTTCTTAATTCAAGTGTCTCTCTG	Hind III
SP058A	NO:321	GACTGGATCCAAATCAATTGGTAGCACAAGATCC	Bam HI
SP058B	NO:322	CAGTGTGACATTAGGAGCCACTGGTCTC	Sal I
SP059A	NO:323	CAGTGGATCCCAAACAGTCAGCTTCAGGAAC	Bam HI
SP059B	NO:324	GACTCTGCAGTTTAATCTTGTCCAGGTGG	Pst I
SP060A	NO:325	GACTGGATCCATTCGATGATGCGGATGAAAAG	Bam HI
SP060B	NO:326	GACTAAGCTTCATTTGTCTTTGGGTATTTTCGCA	Hind III
SP062A	NO:327	CAGTGGATCCGAGAGTCGATCAAAAGTAG	Bam HI
SP062B	NO:328	GTCAGTGCAGTTGCTCGTCTCGAGGTTT	Pst I
SP063A	NO:329	CAGTGGATCCATGGACAACAGGAACTGGGAC	Bam HI
SP063B	NO:330	CAGTAAGCTTATTAGCTTCTGTACCTGTGTTTG	Hind III
SP064A	NO:331	GACTGGATCCCGATGGGCTCAATCCAACCCAGGTCAAGTC	Bam HI
SP064B	NO:332	GACTCTGCAGCATAGCTTTATCCTCTGACATCATCGTATC	Pst I
SP065A	NO:333	GACTGGATCCTTCCAATCAAAAACAGGCAGATGG	Bam HI
SP065B	NO:334	GACTAAGCTTGAGTCCCATAGTCCAAGGCA	Hind III
SP067A	NO:335	AGTCGGATCCTATCACAGGATCGAACGGTAAGACAACC	Bam HI
SP067B	NO:336	ACTGGTGCAGTTCTTTTAACTCCGCTACTGTGTC	Sal I

Table 3

Primer		<i>S. pneumoniae</i> ORF Cloning Primers	
Name	SEQ ID	Sequence	RE
SP068A	NO:337	CAGTGGATCCAAGTTCATCGAAGATGGTTGGGAAGTCC	Bam HI
SP068B	NO:338	GATCGTCGACCCGCTCCCACATGCTCAACCTT	Sal I
SP069A	NO:339	TGACGGATCCATCGCTAGCTAGTGAAATGCAAGAAAAG	Bam HI
SP069B	NO:340	TGACAAGCTTATTCGTTTTTGAAGTAGTTGCTTTTCGT	Hind III
SP070A	NO:341	GACTGGATCCGCACCAGATGGGGCACAAGGTTCAAGG	Bam HI
SP070B	NO:342	TGACAAGCTTAACTTGTAACGAACAGTTCAATCTG	Hind III
SP071A	NO:343	GACTAGATCTTTTTAACCCAAGTGTGGTACTTTCC	Bgl II
SP071B	NO:344	TGACAAGCTTGTTAGGTGTTACATTTTGACCGTC	Hind III
SP072A	NO:345	ACTGAGATCTTTTTAACCCAAGTGTGGTACTTTC	Bgl II
SP072B	NO:346	GACTAAGCTTTCTACGATAACGATCATTTTCTTTACC	Hind III
SP073A	NO:347	GACTGTCGACTCGTAGATATTTAAGTCTAAGTGAAGCG	Sal I
SP073B	NO:348	AGTCAAGCTTGTTAGGTGTTACATTTTGCAAGTC	Hind III
SP074A	NO:349	GACTGGATCCCTTTGGTTTTGAAGGAAGTAAG	Bam HI
SP074B	NO:350	TGACCTGCAGACGATTTTTGAAAAATGGAGGTGTATC	Pst I
SP075A	NO:351	CAGTGGATCCCTACTACCTCTCGAGAGAAAAG	Bam HI
SP075B	NO:352	ACTGAAGCTTTTCGCTTTTTTACTCGTTTGACA	Hind III
SP076A	NO:353	CAGTGGATCCTAAGGTCAAAAGTCAGACCGCTAAGAAAGTGC	Bam HI
SP076B	NO:354	CAGTAAGCTTTAGGGTATCCAAATACTGGTTGTTGATG	Hind III
SP077A	NO:355	TGACAGATCTTGACGGGTCTCAGGATCAGACTCAGG	Bgl II
SP077B	NO:356	TGACAAGCTTCAAAGACATCCACCTCTTGACCTTTG	Hind III
SP078A	NO:357	GACTGGATCCTAGAGGCTTTGCCAAATGGTGGGAAGGG	Bam HI
SP078B	NO:358	GTCAGTCGACTTGTTGTAACACTTTTCGAGGTTTGGTACC	Sal I
SP079A	NO:359	CAGTGGATCCTCAAAAAGAGAAGGAAAAGTGG	Bam HI
SP079B	NO:360	CAGTCTGCAGTTTCTTCAACAAACCTTGTTCTTG	Pst I
SP080A	NO:361	CAGTGGATCCACGTTCTATTGAGGACCACTT	Bam HI
SP080B	NO:362	CAGTAAGCTTTTCTTCTCAGTCAATCTTTTCC	Hind III
SP081A	NO:363	GACTGGATCCCGCTCAAAATACCAGAGGTGTTGAG	Bam HI
SP081B	NO:364	GACTAAGCTTAGTACCATGGGTGTGACAGGTTTGAA	Hind III
SP082A	NO:365	CTGAGGATCCAATTGTACAATTAGAAAAAGATAGC	Bam HI
SP082B	NO:366	TGACAAGCTTGCGTTGACTAGGTTCTGCAATGCC	Hind III
SP083A	NO:367	GACTGGATCCTCTGACCAAGCAAAAAGAAGCAGTCAATGA	Bam HI
SP083B	NO:368	TCAGCAGCTGATCATTGACTTTACGATTTGCTCC	Bgl II
SP084A	NO:369	GACTGGATCCGTCGCGCTCTGTCCAGTCCACTTTTTTCAGCG	Bam HI
SP084B	NO:370	TCAGAAGCTTATTTTTTGTTCCTTAATGCGTT	Hind III
SP085A	NO:371	GACTGGATCCGGGACAAATTCAAAAAATAGGCAAGAGG	Bam HI
SP085B	NO:372	GTCAAAGCTTTGGCTCTTTGATTGCCAACAACTG	Hind III
SP086A	NO:373	GACTGGATCCTCGCTACCAGCAACAAAGCGAGCAAAAAGG	Bam HI
SP086B	NO:374	GACTAAGCTTACTTTTTTCTTTTTTCCACACGA	Hind III
SP087A	NO:375	CAGTGGATCCGAACCGACAAGTCGCCCCTATCAAGACT	Bam HI
SP087B	NO:376	CTGAAAGCTTTGAATTCTCTTTCTTTTCAGGCT	Hind III
SP088A	NO:377	TCGAGGATCCGTTGTGCGGCTGGCAATATATCCCGT	Bam HI
SP088B	NO:378	CAGTAAGCTTCCGAACCCATTGCGCCATTATAGTTGAC	Hind III
SP089A	NO:379	AGTCGGATCCGGCCAAATCAGAATGGGTAGAAGAC	Bam HI
SP089B	NO:380	TGACCTGCAGCTTCTCATTGATTTTCATCATCAC	Pst I
SP090A	NO:381	GACTGGATCCATTTGCAGATGATTCTGAAGGATGG	Bam HI
SP090B	NO:382	TCAGCTGCAGCTTAACCCATTACCATTTCTAGTTAAG	Pst I
SP091A	NO:383	GACTGGATCCTGTGCTGCAAATGAAACTGAAGTAGC	Bam HI
SP091B	NO:384	GACTAAGCTTATACCAAACGCTGACATCTACGCG	Hind III
SP092A	NO:385	AGTCAGATCTTACGTCTCAGCCTACTTTTGTAAGAGC	Bgl II
SP092B	NO:386	GACTAAGCTTAACCCATTACCATTTGGCATTGAC	Hind III
SP093A	NO:387	CAGTGGATCCTGGACAGGTGAAAGGTCATGCTACATTTGTG	Bam HI
SP093B	NO:388	GACTAAGCTTCAACCATTTGAGACCTTGCAACAC	Hind III
SP094A	NO:389	GTCAGGATCCGATTGCTCCTTTTGAAGGATTTGAGAGAAACC	Bam HI
SP094B	NO:390	GACTAAGCTTCGATCAAAGATAAGATAAATATATATAAAGT	Hind III
SP095A	NO:391	GACTGGATCCTAGGTCATATGGGACTTTTTTTCTACAACAAAATAGG	Bam HI



Table 3

Primer		<i>S. pneumoniae</i> ORF Cloning Primers	
Name	SEQ ID	Sequence	RE
SP095B	NO:392	TGACAAGCTTATCTATCAGCTCATTTAATCGTTTTTG	Hind III
SP096A	NO:393	CTGAGGATCCCAACGTTGAGAATTATTTGCGAATG	Bam HI
SP096B	NO:394	TGACAAGCTTGAGTCTACAAAAGTAATGTAC	Hind III
SP097A	NO:395	GTCAGGATCCCTACTATCAATCAAGTTCTTCAGCC	Bam HI
SP097B	NO:396	TGACAAGCTTGACTGAGGCTTGACCAGATTGAAAAG	Hind III
SP098A	NO:397	GACTGGATCCGACAAAAACATTAACCGCTCTGAGG	Bam HI
SP098B	NO:398	GACTAAGCTTAGCACGAACGTGTGACGCTGGTTCC	Hind III
SP099A	NO:399	GACTGGATCCTTCTCAGGAGACCTTTAAAAATATC	Bam HI
SP099B	NO:400	GACTAAGCTTGTTGGCCATCTTGACATACC	Hind III
SP100A	NO:401	GACTGGATCCAGTAAATGCGCAATCAAATTC	Bam HI
SP100B	NO:402	AGTCCTGCAGGTATTTAGCCCAATAATCTATAAAGCT	Pst I
SP101A	NO:403	CAGTGGATCCTTACCGCGTTCATCAAGATGTC	Bam HI
SP101B	NO:404	GACTAAGCTTGCCAGATGTTGAAAAGAGAGTG	Hind III
SP102A	NO:405	GACTGGATCCGTGGATGGGCTTTAACTATCTTCGTATTCCG	Bam HI
SP102B	NO:406	AGTCAAGCTTGCTAGTCTTCACTTTCCCTTTCC	Hind III
SP103A	NO:407	GACTGTCGACACTAAACCAGCATCGTTCGCAGGA	Sal I
SP103B	NO:408	CTGACTGCAGCTTCTTGAAGAAATAATGATTGTGG	Pst I
SP105A	NO:409	CAGTGGATCCTGACTACCTTGAAATCCCACTT	Bam HI
SP105B	NO:410	CAGTAAGCTTTTTTTTAAAGTTGTAGAATGATTTCAATC	Hind III
SP106A	NO:411	CAGTGTGCACTCGTATCTTTTTTTGGAGCAATGTT	Sal I
SP106B	NO:412	GACTAAGCTTAAATGTTCCGATACGGGTGATTG	Hind III
SP107A	NO:413	CAGTGGATCCGGACTCTCTCAAAGATGTGAAAG	Bam HI
SP107B	NO:414	GACTAAGCTTCTTGAGTTTGTCAAGGATTGCTTT	Hind III
SP108A	NO:415	CAGTGGATCCCAAGAAATCCTATCATCTCTTCCAGAAG	Bam HI
SP108B	NO:416	GACTAAGCTTTTTCAGAACTAAAGCCGCAGCTT	Hind III
SP109A	NO:417	GACTGGATCCACGAAATGCAGGGCAGACAG	Bam HI
SP109B	NO:418	CAGTAAGCTTATCAACATAATCTAGTAAATAAGCGT	Hind III
SP110A	NO:419	CAGTGGATCCTGTATAGTTTTTACGCGTTGTTCTTC	Bam HI
SP110B	NO:420	GTCAAAGCTTTGATAGAGTGTCAATCTTCTTTAG	Hind III
SP111A	NO:421	GACTGGATCCGTGTGTCGAGCATATTCTGAAG	Bam HI
SP111B	NO:422	CAGTAAGCTTACTTTTTACCAATTTCTTGTCTGCATC	Hind III
SP112A	NO:423	GACTGTCGACGTGTTTGGATAGCATTCAGAATCAGACG	Sal I
SP112B	NO:424	CAGTAAGCTTCGGAAGTAAAGACAATTTTTTCC	Hind III
SP113A	NO:425	CAGTGGATCCGTGCCCTAGATAGTATTACTCAAAC	Bam HI
SP113B	NO:426	GACTAAGCTTTTTTGCTTATTTCTCTCAATTTTTTC	Hind III
SP114A	NO:427	CAGTGGATCCCATTCAGAAGCAGACCTATCAAATC	Bam HI
SP114B	NO:428	ACTGAAGCTTATGTAATTTTTTAGATTTTTCAATATTTTTAG	Hind III
SP115A	NO:429	AGTCGGATCCTAAGGCTGATAATCGTGTTCAAATG	Bam HI
SP115B	NO:430	GACTAAGCTTAAATTAGATAGACGTTGAGT	Hind III
SP117A	NO:431	AGTCGGATCCCTGTGGCAATCAGTCAGCTGCTTCC	Bam HI
SP117B	NO:432	GACTGTCGACTTTAATCTTGTCCAGGTGGTTAATTTGCC	Sal I
SP118A	NO:433	ACTGGTCGACTTGTCAACAACAACATGCTACTTCTGAG	Sal I
SP118B	NO:434	GACTCTGCAGAAGTTTAACCCACTTATCATTATCC	Pst I
SP119A	NO:435	ACTGGGATCCTTGTTCAGGCAAGTCCGTGACTAGTGAAC	Bam HI
SP119B	NO:436	GACTAAGCTTGGCTAATTCCTTCAAAGTTTGCA	Hind III
SP120A	NO:437	AGTCGGATCCCTCGCAAATGAAAAGGCGGCAGTTAGCC	Bam HI
SP120B	NO:438	GACTAAGCTTGTAATAAGCGTACCTTTTTCTTCC	Hind III
SP121A	NO:439	TCAGGGATCCTTGTCACTCAGGTTCTAATGGTTCTCAG	Bam HI
SP121B	NO:440	AGTCAAGCTTGGCATTGGCGTCGCCGTCCTTC	Hind III
SP122A	NO:441	GACTGGATCCGGAACTTCACAGGATTTTAAAGAGAAG	Bam HI
SP122B	NO:442	GACTGTCGACAATCAATCCTTCTTCTGCACTTCT	Sal I
SP123A	NO:443	CAGTGGATCCTGTGGTCAAGTTGAGACTCCTCAATC	Bam HI
SP123B	NO:444	GACTAAGCTTTTCTTCAAATTTATTATCAGC	Hind III
SP124A	NO:445	AGTCGGATCCAACACCTGTATATAAAGTTACAGCAATCG	Bam HI
SP124B	NO:446	GACTGTCGACTACTTGACCGAATGCGTCAATGTACG	Sal I

Table 3

<i>S. pneumoniae</i> ORF Cloning Primers			
Primer			
<u>Name</u>	<u>SEQ ID</u>	<u>Sequence</u>	<u>RE</u>
SP125A	NO:447	CTGAGGATCCATTAGACAGATTAATTGAAATCGG	<i>Bam</i> HI
SP125B	NO:448	GACTGTCGACTTTTAAAGATTGAAGTTTAAAGCT	<i>Sal</i> I
SP126A	NO:449	TGACGGATCCTAAGACAGATGAACGGAGCAAGGTG	<i>Bam</i> HI
SP126B	NO:450	CTGAAAGCTTTAAGGCTTCCTCAATGAGTTTGTCT	<i>Hind</i> III
SP127A	NO:451	GACTGGATCCCTGTGAGAATCAAGCTACACCCA	<i>Bam</i> HI
SP127B	NO:452	CTGAAAGCTTTTGTAAGTGAAGATTGATCTGGGAG	<i>Hind</i> III